Review

Mutations Induced by Oxidized DNA Precursors and Their Prevention by Nucleotide Pool Sanitization Enzymes

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Base oxidation occurs in the cellular nucleotide pool as well as in DNA, and the oxidized DNA precursors induce mutagenic events. 8-Hydroxy-dGTP (8-OH-dGTP) and 2-hydroxy-dATP (2-OH-dATP) have been identified as the major products of *in vitro* oxidation reactions with Fe^{2+} . The mutagenic potentials of many oxidized DNA precursors have been examined in various experimental systems. Accumulating evidence indicates the importance of 8-OH-dGTP and 2-OH-dATP in the mutation process. In addition, nucleotide pool sanitization enzymes, such as MutT and MTH1, have been identified as a defense against the mutagenesis induced by these oxidized DNA precursors. In this review, the mutagenicities of 8-OH-dGTP and 2-OH-dATP and the functions of MutT-type nucleotide pool sanitization enzymes will be summarized.

Key words: oxidized DNA precursor, 8-hydroxy-dGTP, 2-hydroxy-dATP, nucleotide pool sanitization enzyme

Introduction

Reactive oxygen species (ROS) are produced endogenously through normal cellular metabolism, and exogenously by various environmental mutagens and dietary factors (1-4). Various kinds of DNA lesions are generated by ROS, and the mutational properties of many DNA lesions have been examined (5). The formation of DNA lesions is recognized as being related to mutagenesis, carcinogenesis, neurodegeneration, and aging (1,2,6).

Base oxidation occurs in the cellular nucleotide pool as well as in DNA, and the oxidized deoxyribonucleotides (DNA precursors) induce mutagenic events (5,7). Previously, the *in vitro* oxidation products of dATP and dGTP were analyzed, and the amounts of oxidatively damaged bases were quantitated. DNA precursors were more susceptible to the oxidation than DNA. The amount of 8-hydroxyguanine (8-OH-Gua) formed upon the treatment of dGTP with Fe^{2+} -EDTA is ~9-fold more abundant than that in DNA with the same treatment (8). Likewise, 2-hydroxyadenine (2-OH-Ade) is formed ~67-fold more efficiently upon dATP oxidation than upon DNA oxidation. Intracellular DNA is bound

with proteins, such as histones, and its reactivity would be low, due to the protection by the proteins. Thus, the oxidation of DNA precursors in cells may play more important roles in ROS-induced mutagenesis than previously speculated. The accumulation of 8-OH-Gua by the incorporation of oxidized dGTP from the nucleotide pool contributes almost equally to the direct oxidation of G bases in DNA (9). Nunoshita *et al.* reported that the mutations found in an *Escherichia coli* strain lacking superoxide dismutases and a repressor for iron-uptake systems were A:T→C:G and G:C→T:A transversions, and concluded that these mutations would be caused by 8-hydroxy-2'-deoxyguanosine 5'-triphosphate (8-OH-dGTP) and 2-hydroxy-2'-deoxyadenosine 5'-triphosphate (2-OH-dATP), respectively, on the basis of various experiments (10). These results suggest that oxidatively damaged DNA precursors are biologically quite important.

Experiments with synthetic deoxyribonucleoside triphosphates provide clear data on the relationship between the damaged structures and the mutational properties. In this review, the author will particularly focus on 8-OH-dGTP and 2-OH-dATP (Fig. 1), which are the major products of *in vitro* oxidation reactions with Fe^{2+} (11). In addition, the nucleotide pool sanitization enzymes that function as a defense against the mutagenesis induced by oxidatively damaged DNA precursors will be discussed.

Incorporation by Purified DNA Polymerases

The incorporation of 8-OH-dGTP by various purified DNA polymerases (pols) has been examined *in vitro*, using synthetic oligodeoxyribonucleotides as primed templates. The oxidized form of dGTP is incorporated opposite C and A in the templates (Table 1). The ratio of the incorporation of 8-OH-dGTP opposite A to that opposite C varies, depending upon the DNA pol (12-16).

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Fig. 1. Structures of 8-OH-dGTP and 2-OH-dATP. The equilibrium between the keto (left) and hydroxy (right) forms shifts to the keto form for 8-OH-dGTP. The proportion of the two forms of 2-OH-dATP is affected by the environment around the compound (60,61). Ⓟ represents a phosphate group.

Table 1. Mutations induced by 8-OH-dGTP and 2-OH-dATP in various experimental systems

DNA precursors	Experimental system		
	Purified DNA pol	<i>E. coli</i>	Human cell extract
8-OH-dGTP	Misincorporation opposite A	A:T→C:G	A:T→C:G
2-OH-dATP	Misincorporation opposite C and G	G:C→T:A	G:C→A:T G:C→T:A

Shimizu *et al.* have reported that the Y-family DNA pols, including *Sulfolobus acidocaldarius* DNA pol Y1 (Dbh), *S. solfataricus* strain P2 DNA pol Y1 (Dpo4), *E. coli* DNA pol IV (DinB), human DNA pol η (XPV), and pol ι , incorporate 8-OH-dGTP opposite A with quite high frequencies (17–19). 8-OH-dGTP was added to the deoxyribonucleotides used for the gap-filling reaction by the Klenow fragment of *E. coli* DNA pol I, and the synthesized DNA was transfected into *E. coli* (20). A→C transversions were almost exclusively detected. Similar results were obtained when other DNA pols, including the *E. coli* DNA pol III holoenzyme, were used (21,22). The A→C transversions observed in these studies imply that 8-OH-dGTP is incorporated opposite A, and then dCTP is inserted opposite 8-OH-Gua in the DNA (Fig. 2).

The incorporation of 2-OH-dATP by purified DNA pols *in vitro* has also been examined, using synthetic oligodeoxyribonucleotides as primed templates (Table 1). The *E. coli* DNA pol III α subunit incorporates 2-OH-dATP opposite T and G (23). The kinetic parameters of the incorporation indicated that the efficiencies opposite T and G are ~10:1. On the other hand, the mammalian replicative DNA pol, pol α , incorporates 2-

Fig. 2. Proposed model for mutations induced by 8-OH-dGTP and 2-OH-dATP. 2-OH-dATP is misincorporated opposite G and C in a DNA pol-dependent manner. G^{OH}, 8-OH-Gua; A^{OH}, 2-OH-Ade.

OH-dATP opposite T and C with ~4.5:1 efficiencies (8). In contrast to 8-OH-dGTP, the incorporation mode of 2-OH-dATP is DNA pol-dependent. Interestingly, the DNA pol-dependency is observed for mispair formation of 2-OH-Ade in DNA templates (24,25). Shimizu *et al.* reported that the incorporation of 2-OH-dATP opposite T and G occurs with similar efficiencies

by the Y-family DNA pols (17–19). The gap-filling reaction by the *E. coli* DNA pol III holoenzyme was examined, and G→T transversions were found to be induced (22). It seems that 2-OH-dATP is incorporated opposite G, and then dTTP is inserted opposite 2-OH-Ade in the DNA (Fig. 2).

The misincorporation of other damaged DNA precursors has been studied using various purified DNA pols, as described in detail in another review (5).

Mutations Induced in *E. coli*

The mutagenicity of 8-OH-dGTP and 2-OH-dATP *in vivo* was examined by the direct incorporation of damaged DNA precursors, by treating *E. coli* cells with CaCl_2 and isolating the mutants of a target gene (26). The treatments with 8-OH-dGTP and 2-OH-dATP increased the frequency of substitution mutations, in contrast to the treatments with dGTP and dATP. This result indicates that damaged DNA precursors act as mutagens in cells. The substitution mutation found most frequently in the 8-OH-dGTP-induced mutants is an A:T→C:G transversion (Table 1). This type of mutation comprised 90% of the substitution mutations detected, in the case of the *lacI* gene (26). The substitution mutation found most frequently in the 2-OH-dATP-induced mutants was a G:C→T:A transversion (84% of the total substitution mutations detected in the *lacI* gene). As described in the Introduction section, Nunoshiba *et al.* reported that the mutations found in an *E. coli* strain lacking superoxide dismutases and a repressor for iron-uptake systems were A:T→C:G and G:C→T:A transversions, and concluded that these mutations would be caused by 8-OH-dGTP and 2-OH-dATP, respectively (10). Thus, these results suggest that the 8-OH-dGTP and 2-OH-dATP formed in the nucleotide pool contribute to the mutations induced by ROS. The A:T→C:G and G:C→T:A transversions observed for 8-OH-dGTP and 2-OH-dATP, respectively, would be initiated by their misincorporations opposite A and G (Fig. 2).

When 5-hydroxy-2'-deoxycytidine 5'-triphosphate and 5-formyl-2'-deoxyuridine 5'-triphosphate, oxidized dCTP and dTTP, respectively, were introduced directly into *E. coli*, these deoxyribonucleotides induced *lacI*[−] and *lacO*^c mutants as efficiently as 8-OH-dGTP (27). 5-Hydroxy-2'-deoxycytidine 5'-triphosphate and 5-formyl-2'-deoxyuridine 5'-triphosphate induced A:T→C:G, A:T→G:C, and G:C→T:A mutations, and G:C→A:T, A:T→G:C, and G:C→T:A mutations, respectively. In contrast, 8-hydroxy-2'-deoxyadenosine 5'-triphosphate (8-OH-dATP), which is formed by ionizing radiation, caused little, if any, mutagenesis (28). The mutagenicity of 8-OH-dATP is at least one order of magnitude lower than that of 2-OH-dATP in *E. coli*. In addition, formamidopyrimidine-dGTP (Fapy-dGTP,

*N*⁶-(2-deoxy- α , β -D-ribofuranosyl)-2,6-diamino-5-formamido-4-hydroxypyrimidine 5'-triphosphate), another oxidation product of dGTP, is nonmutagenic in *E. coli* (Kamiya *et al.*, unpublished results).

Mutations Induced in *in vitro* DNA Replication with Extracts of Human Cells

The effects of 8-OH-dGTP on the fidelity of replication conducted by a HeLa cell extract have been reported (Table 1) (21). The replicated DNA was transfected into *E. coli*, and A:T→C:G mutations were induced. This induction of A:T→C:G mutations by 8-OH-dGTP was dependent on the MutY activity: The frequency of these mutations decreased when *mutY* strains were used as host cells (21). The mutagenicity of 2-OH-dATP was also examined, using an *in vitro* replication system with a HeLa extract. 2-OH-dATP induced substitution and deletion mutations (29). Among the substitutions, a G:C→A:T transition including a tandem (CC→TT) mutation was mainly induced. This result is in contrast to the finding that G:C→T:A transversions are induced by 2-OH-dATP in *E. coli* cells (Table 1). This discrepancy can be explained by the DNA pol-dependent misincorporation mode of 2-OH-dATP (8,23), which is incorporated opposite C in the extract, as by purified DNA pol α (Fig. 2).

Recently, we observed that 2-OH-dATP synergistically enhanced the mutagenicity of 8-OH-dGTP in the reaction (30). 2-OH-dATP suppressed the hydrolysis of 8-OH-dGTP, suggesting that the inhibition of an MTH1(-like) protein played a major role in the enhancement (see below). These results highlight the importance of the specific hydrolysis of 8-OH-dGTP for the suppression of its induced mutation.

Mutations Induced in Live Mammalian Cells

To examine the mutagenicity of 8-OH-dGTP in mammalian cells, 8-OH-dGTP and shuttle plasmid DNA containing a target gene were co-introduced by means of cationic liposomes into simian COS-7 cells. The DNAs replicated in the cells were recovered and then transfected into *E. coli*. The 8-OH-dGTP induced A:T→C:G substitution mutations in the COS-7 cells (31). This result agrees with previous observations that DNA pols misincorporated 8-OH-dGTP opposite A *in vitro*, and that this oxidized dGTP induced A:T→C:G transversions in *E. coli*. As described above, the observed A:T→C:G transversions observed can be explained by the incorporation of 8-OH-dGTP opposite A and the insertion of dCTP opposite 8-OH-Gua in DNA during the next round of replication (Fig. 2). These results constitute the first direct evidence to show that 8-OH-dGTP actually induces mutations in living mammalian cells.

Roles of Y-family DNA Polymerases in Mutation Induction

As described above, some Y-family DNA pols incorporate damaged DNA precursors opposite incorrect bases with high frequencies *in vitro* (17–19). To reveal the roles of Y-family DNA pols in the mutagenesis induced by oxidatively damaged DNA precursors, 2-OH-dATP and 8-OH-dGTP were introduced into *E. coli* strains deficient in the Y-family DNA pols, DNA pol IV (DinB) and DNA pol V (UmuD'₂C) (32). The mutation induced by 2-OH-dATP, but not that induced by 8-OH-dGTP, occurred less frequently in the *dinB* strain than in the wild-type (wt) strain, suggesting the involvement of DNA pol IV in the mutagenesis by 2-OH-dATP. The expression of DNA pol IV from a plasmid enhanced the mutagenesis by 2-OH-dATP in the *dinB* strain. This enhancement was dependent on the DNA pol activity, since the expression of a mutant DNA pol IV lacking the pol activity did not increase the mutations induced by 2-OH-dATP. As revealed by the *in vitro* DNA synthesis experiment (18), DNA pol IV seems to facilitate the mutagenicity of 2-OH-dATP through the misincorporation of 2-OH-dATP. Moreover, DNA pol IV may be involved in the extension from the 3'-terminal 2-OH-Ade residue. Alternatively, DNA pol IV may function in the fixation of mutations, by the incorporation of dTTP opposite 2-OH-Ade during the second round of replication and/or the extension from the 3'-terminal T residue incorporated opposite 2-OH-Ade (Fig. 2). In contrast, both 8-OH-dGTP and 2-OH-dATP caused mutations more efficiently in the *umuDC* strain than in the wt strain, suggesting that the *umuDC* gene products suppressed the mutagenesis by these oxidized DNA precursors (32). The DNA pol activity was not required for the suppressive effects, since the expression of the *umuDC* gene products lacking the DNA pol activity also suppressed the mutagenesis. These results suggest that *E. coli* DNA pol IV is involved in the mutagenesis by 2-OH-dATP, and that the *umuDC* gene products play suppressive role(s) in the mutagenesis by both 8-OH-dGTP and 2-OH-dATP.

The mutagenicity of 8-OH-dGTP in an *in vitro* replication system with an extract of XP-V cells lacking a Y-family DNA pol, pol η , was similar to that with a HeLa extract (30). This result suggests that the contribution of this highly error-prone DNA pol is low in this experimental system.

In contrast, however, the incorporation of 8-OH-dGTP by DNA pol η may induce mutations in living human cells, since the mutagenicity of 8-OH-dGTP directly incorporated into cells was reduced significantly when the expression of DNA pol η was disabled by siRNA (Satou *et al.*, unpublished results).

Nucleotide Pool Sanitization Enzymes in *E. coli*

MutT is the gene product of the *E. coli* mutator gene, *mutT* (*nudA*). Disruption of the *mutT* gene led to an increased frequency of A:T→C:G mutations (9,33). In 1992, Maki and Sekiguchi found that the MutT protein specifically hydrolyzed 8-OH-dGTP to form the corresponding monophosphate (Table 2) (12). This report highlighted the role of 'nucleotide pool sanitization' to prevent the mutagenesis induced by damaged DNA precursors. The facts that the accumulation of 8-OH-Gua by the incorporation of 8-OH-dGTP contributed almost equally to the direct oxidation of G bases in DNA (9), and that the A:T→C:G transversions occurred with high frequency in the *mutT* strain, indicate the importance of nucleotide pool sanitization as the first defense system against damaged DNA precursors. In agreement with this interpretation, exogenous 8-OH-dGTP induced mutations more efficiently in an *mutT* strain than in the wt strain (34). This result constitutes the first direct evidence that the MutT protein suppresses mutations by 8-OH-dGTP *in vivo*. Recently, 8-hydroxy 2'-deoxyguanosine 5'-diphosphate (8-OH-dGDP) was found to be another substrate for the MutT protein (35). Although the biological significance of this enzymatic activity is unknown, the authors of the paper suggested that 8-OH-dGTP and 8-OH-dGDP are interconvertible by the actions of nucleoside diphosphate kinase and nucleoside triphosphatase, which do not discriminate these damaged deoxyribonucleotides from the normal ones (36).

The *E. coli* Orf135 (NudG) protein is another MutT-type enzyme and contains the 'MutT signature/phosphohydrolase module/Nudix box' (37–39). This protein catalyzes the hydrolysis of 2-OH-dATP, and somewhat less efficiently, 8-OH-dGTP (Table 2) (40). Exogenous 2-OH-dATP induced mutations more efficiently in an *orf135* strain than in the wt strain (41), suggesting that the Orf135 protein catalyzes the hydrolysis of 2-OH-dATP *in vivo*. On the other hand, 8-OH-dGTP induced mutations similarly in the *orf135* and wt

Table 2. MutT-type nucleotide pool sanitization enzymes

Proteins	Candidate substrates
<i>E. coli</i>	
MutT	8-OH-dGTP, 8-OH-dGDP
Orf17	8-OH-dATP, 2-OH-dATP, 8-OH-dGTP 8-OH-dADP, 2-OH-dADP, 8-OH-dGDP
Orf135	2-OH-dATP, 8-OH-dGTP
<i>S. cerevisiae</i>	
PCD1	8-OH-dGTP, 2-OH-dATP
Mammalian cells	
MTH1	8-OH-dGTP, 2-OH-dATP, 8-OH-dATP
MTH2	8-OH-dGTP
NUDT5	8-OH-dGDP

strains. This result suggests that the MutT protein, rather than the Orf135 protein, primarily degrades 8-OH-dGTP in *E. coli*. The spontaneous and H₂O₂-induced mutant frequencies were two- to three-fold higher in the *orf135* strain than in the wt strain (41). Various mutations involving a G:C→T:A transversion, the same type of mutation induced by 2-OH-dATP, occurred more frequently in the *orf135* strain than in the wt strain.

Orf135 proteins with substitutions at amino acid residues in the MutT signature sequence were generated, and the *in vitro* enzymatic activities were measured (42). In addition, the mutant Orf135 proteins were each expressed in the *orf135* strain, and the mutation frequency upon H₂O₂ treatment was examined (43). The *in vivo* mutation suppression abilities and the *in vitro* enzymatic activities were then compared. The expression of the enzymatically active Orf135 mutants in the *orf135* strain tended to reduce the mutation frequency induced by H₂O₂ (43). This result suggests the importance of the phosphohydrolase activity in the suppression of mutations by the Orf135 protein. Mutant Orf135 proteins with substitutions at amino acid residues that are putatively involved in the base recognition of substrates were generated, and the *in vitro* enzymatic activities and the *in vivo* mutation suppression abilities were compared (44). A statistically significant correlation was found only between the H₂O₂-induced mutation frequencies and the 2-OH-dATPase activity (albeit partially). The results of these experiments suggest that the 2-OH-dATPase activity of the Orf135 protein contributes to the suppression of the ROS-induced mutagenesis derived from oxidized deoxyribonucleotide(s) in *E. coli* cells.

Another MutT-type enzyme, Orf17 (NudB), catalyzes the hydrolysis of 8-OH-dATP and, less efficiently, 8-OH-dGTP and 2-OH-dATP (Table 2) (45). The Orf17 protein also catalyzes the hydrolysis of their diphosphate derivatives. Since the destruction of the *orf17* gene results in severe growth inhibition, plasmid DNA containing the gene in the antisense orientation, under an inducible promoter, was introduced into *E. coli* (46). Although a slightly increased mutation frequency was observed for spontaneous mutations, no clear phenotype was observed for H₂O₂- and X-ray-induced mutations. Moreover, the introduction of 8-OH-dATP into the antisense-expressing cells did not induce mutations (Hori *et al.*, unpublished results). Although it is possible that 8-OH-dATP itself is not mutagenic (28), the Orf17 protein may not contribute to the suppression of ROS-induced mutations. On the other hand, the overexpression of the Orf17 protein in *mutT* cells decreased the mutation frequency (46). This result suggests that the Orf17 protein could act as a backup protein for the MutT protein.

The RibA protein (GTP cyclohydrolase II), which is involved in riboflavin synthesis, catalyzes the hydrolysis of 8-OH-dGTP *in vitro* (47). The *ribA* gene reduced the increased level of mutation frequency in the *mutT* strain to almost the normal level, when the gene product was overproduced. In the *mutT* background, the *ribA* cells showed higher spontaneous mutation frequencies, as compared with that of *ribA*⁺ cells. Thus, GTP cyclohydrolase II has the potential to protect genetic material from the untoward effects of ROS. Although the RibA protein is not a member of the Nudix superfamily, it could act as a backup protein for the MutT protein.

Nucleotide Pool Sanitization Enzyme in Yeast

Nunoshiba *et al.* found that the *Saccharomyces cerevisiae* PCD1 (YLR151c) protein, a member of the Nudix superfamily, catalyzes the hydrolysis of 8-OH-dGTP and 2-OH-dATP (Table 2) (48). Expression of PCD1 suppressed the A:T→C:G transversion in the *E. coli mutT* strain. The disruption of the PCD1 gene in yeast caused a 14-fold increase in the frequency of spontaneous mutations, as compared to the wt strain. Thus, PCD1 has the ability to prevent spontaneous mutagenesis *via* sanitization of the nucleotide pool, and therefore, it may be the functional homologue of *E. coli* MutT in *S. cerevisiae*.

Nucleotide Pool Sanitization Enzymes in Mammalian Cells

In mammalian cells, the MTH1 protein, a member of the MutT-type enzymes containing the MutT signature/phosphohydrolase module/Nudix box, hydrolyzes 8-OH-dGTP (Table 2) (49,50). When the cDNA for human MTH1 was expressed in *E. coli mutT* mutant cells, the elevated level of spontaneous A:T→C:G mutations reverted to the normal level (51). This protein also catalyzes the hydrolysis of 2-OH-dATP and 8-OH-dATP *in vitro* (52). Tsuzuki *et al.* reported that a greater number of tumors were spontaneously formed in the lungs, livers, and stomachs of MTH1-deficient mice, as compared with wt mice (53). Thus, the accumulation of 8-OH-dGTP and 2-OH-dATP in the nucleotide pool might induce mutations in tumor-related genes, thus enhancing tumorigenesis. Nakabeppu and collaborators generated mutant MTH1 proteins that lacked either the 8-OH-dGTPase or 2-OH-dATPase activity (54). Using these mutant MTH1 proteins, they revealed that the H₂O₂-induced cell dysfunction and death observed in MTH1-null mouse embryo fibroblasts were effectively suppressed by the expression of wt human MTH1, and that they were only partially suppressed by the expression of a mutant MTH1, defective in either the 8-OH-dGTPase or 2-OH-dATPase activity (55). 2-OH-dATP synergistically enhanced the mutagenicity of 8-OH-dGTP in *in vitro* DNA replication with an extract of hu-

man cells (30). 2-OH-dATP suppressed the hydrolysis of 8-OH-dGTP, suggesting that the inhibition of the MTH1(-like) protein plays a major role in the enhancement. These results highlight the importance of the specific hydrolysis of 8-OH-dGTP for the suppression of its induced mutation.

Another member of the mammalian MutT-type enzymes is the MTH2 protein. The protein obtained by the expression of the cloned mouse MTH2 cDNA catalyzes the hydrolysis of 8-OH-dGTP (56). Expression of the cDNA encoding MTH2 significantly reduced the elevated level of spontaneous mutation frequency in *E. coli mutT* cells. Thus, MTH2 could act as a backup protein for the MTH1 protein.

The human Nudix type 5 (NUDT5) protein catalyzes the hydrolysis of 8-OH-dGDP to the monophosphate derivative (57). On the other hand, the hydrolysis of 8-OH-dGTP is catalyzed by the NUDT5 protein only at very low levels. However, the increased frequency of spontaneous mutations in *E. coli mutT* cells is decreased to normal levels when NUDT5 is expressed. This result suggests that the elimination of the damaged deoxyribonucleoside diphosphate from the nucleotide pool is important for the prevention of mutations. As described above, 8-OH-dGTP and 8-OH-dGDP are interconvertible in cells, and the removal of either deoxyribonucleotide may contribute to mutation prevention to similar degrees.

The reports described above indicate that the nucleotide pool sanitization enzymes contribute to the suppression of mutations, and that damaged DNA precursors are potentially endogenous mutagens. The fact that *E. coli*, yeast, and mammalian cells possess enzyme(s) with 2-OH-dATPase activity indicates that 2-OH-dATP is an important oxidized DNA precursor.

Roles of DNA Repair Proteins

As described above, the nucleotide pool sanitization enzymes act as the first defense against the mutagenesis by damaged DNA precursors. Alternatively, DNA repair enzymes may prevent the mutagenesis by the oxidized deoxyribonucleotides after their incorporation into DNA. We recently examined this possibility, using *E. coli* strains deficient in a gene related to DNA repair. Our data suggested that endonuclease III (Nth) functions as the second defense against 8-OH-dGTP (58). In contrast, UvrA and UvrB enhanced the mutations induced by oxidized deoxyribonucleotides by unknown mechanism(s) (59). The roles of DNA repair proteins in the mutagenesis by damaged DNA precursors should be elucidated in detail.

Conclusions

Damaged DNA precursors formed in the nucleotide pool by the action of ROS contribute to mutation induc-

tion. The nucleotide pool sanitization enzymes function as the first defense against the damaged DNA precursors. A thorough understanding the roles of specialized DNA pols and DNA repair proteins is necessary to gain a clear view of the effects of oxidized DNA precursors.

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