Measurement and Meaning of Oxidatively Modified DNA Lesions in Urine

Marcus S. Cooke,1,2 Ryszard Olinski,3 and Steffen Loft4 members of the European Standards Committee on Urinary (DNA) Lesion Analysis (ESCUA)

1Radiation and Oxidative Stress Section, Department of Cancer Studies and Molecular Medicine and 2Department of Genetics, University of Leicester, Leicester Royal Infirmary, University Hospitals of Leicester NHS Trust, Leicester, United Kingdom; 3Department of Clinical Biochemistry, Collegium Medicum, Nicolaus Copernicus University, Bydgoszcz, Poland; and 4Environmental and Occupational Health, Institute of Public Health, University of Copenhagen, Copenhagen, Denmark

Abstract

Background: Oxidatively generated damage to DNA has been implicated in the pathogenesis of a wide variety of diseases. The noninvasive assessment of such damage, i.e., in urine, and application to large-scale human studies are vital to understanding this role and devising intervention strategies.

Methods: We have reviewed the literature to establish the status quo with regard to the methods and meaning of measuring DNA oxidation products in urine.

Results: Most of the literature focus upon 8-oxo-7,8-dihydro-2′-deoxyguanosine (8-oxoG), and whereas a large number of these reports concern clinical conditions, there remains (a) lack of consensus between methods, (b) possible contribution from diet and/or cell death, (c) no definitive DNA repair source of urinary 2′-deoxyribonucleoside lesions, and (d) no reference ranges for healthy or diseased individuals.

Conclusions: The origin of 8-oxoG is not identified; however, recent cell culture studies suggest that the action of Nudix hydrolase(s) on oxidative modification of the nucleotide pool is a likely candidate for the 8-oxoG found in urine and, potentially, of other oxidized 2′-deoxyribonucleoside lesions. Literature reports suggest that diet and cell death have minimal, if any, influence upon urinary levels of 8-oxoG and 8-oxo-7,8-dihydroguanine, although this should be assessed on a lesion-by-lesion basis. Broadly speaking, there is consensus between chromatographic techniques; however, ELISA approaches continue to overestimate 8-oxoG levels and is not sufficiently specific for accurate quantification. With increasing numbers of lesions being studied, it is vital that these fundamental issues are addressed. We report the formation of the European Standards Committee on Urinary (DNA) Lesion Analysis whose primary goal is to achieve consensus between methods and establish reference ranges in health and disease. (Cancer Epidemiol Biomarkers Prev 2008; 17(1):3–14)

Introduction

Reactive oxygen species are produced continually from normal cellular metabolism but may be produced in excess, leading to oxidative stress, after exposure to xenobiotics, radiation, etc. A consequence of reactive oxygen species production is their interaction with, and subsequent modification of, cellular biomolecules, in particular, DNA, lipids, and proteins. Damage to DNA is of particular importance, in part due to the possibility of inheritable sequence alterations (mutations), although it should not be forgotten that reactive oxygen species—induced damage to DNA may have nonmutational effects, such as the promotion of microsatellite instability and acceleration of telomere shortening (1). Over 30 base modifications, such as 8-oxo-7,8-dihydroguanine (8-oxoGua) and its corresponding deoxyribonucleoside equivalent, 8-oxo-7,8-dihydro-2′-deoxyguanosine (8-oxoDG; Fig. 1), have been described (2) with >70 DNA lesions identified (3). Furthermore, reactive intermediates arising from the interaction of reactive oxygen species with lipids and proteins may also modify DNA, leading to adducts, such as 1,N6-etheno-2′-deoxyadenosine (εdA), 3-(2-deoxy-β-D-erythro-pentofuranosyl)pyrimido[1,2-a]purin-10(3H)-one, and DNA-protein cross-links (Fig. 1; ref. 4). A great deal of evidence exists, which suggests that this plethora of lesions, arising from oxidative stress, may have an important role in the etiology and/or pathogenesis of many diseases (reviewed in ref. 5), such as cancer and aging.

To fully understand the extent to which such DNA lesions are involved in disease, methods for their analysis are essential. Numerous approaches have been applied to
the study of oxidatively damaged DNA, including gas chromatography with mass spectrometry (GC/MS; ref. 6), high performance liquid chromatography with electrochemical detection (HPLC ECD; ref. 7), HPLC with single (8) or tandem (9) mass spectrometry, 32P postlabeling (10), immunoassay (11, 12), alkaline elution (13), and Comet assay (14), plus other methods based upon the nicking of DNA at oxidized bases by means of repair enzymes (15, 16). However, after the publication of the findings from the European Standards Committee on Oxidative DNA Damage (17), a number of these techniques have fallen from favor (reviewed in ref. 18).

Assessment of damage to DNA by methods requiring invasive procedures, e.g., blood samples or tissue biopsy, imposes severe limitations in large-scale human studies, requiring staff with specialist training and reducing the likelihood of consent. Furthermore, the possibility of adventitious oxidation during sample storage and DNA extraction has not been entirely ruled out, but methods have been developed to minimize the risk (16). Examining the products of oxidatively generated damage to DNA in extracellular matrices offers a means by which oxidative stress may be assessed noninvasively or minimally invasively and circumvents extraction and the resultant artifacts.

Methods of Analysis

For the most part, methods that have been applied to the analysis of oxidatively damaged DNA lesions in urine are either chromatographic (principally, HPLC MS/MS; liquid chromatography prepurification before GC/MS, HPLC GC/MS, HPLC ECD, GC/MS) or immunoassay. The majority of assays focus upon 8-oxodG as the analyte of choice, although the following have been reported to be present in urine, in many cases with their corresponding deoxyribonucleoside (Table 1).

Chromatographic Techniques. Urine is a very complex matrix, and therefore, the common challenge for all chromatographic techniques has been to clean-up the urine sufficiently to simplify analysis, which very often also extends instrument life. At its simplest, column switching has meant that, after chromatographic separation of the urine’s constituents, only the fraction containing the compound of interest (e.g., 8-oxodG) is applied to the final separation column and detector, either ECD (19, 20) or MS (21); the remainder is diverted to waste.

Using HPLC, thymine glycol, which was a “significant marker” of oxidative stress (22), was one of the first reactive oxygen species-induced DNA lesions to be studied in urine. The methodology required the use of boronate, solid phase extraction columns, and semipreparative HPLC to isolate thymine glycol and thymidine glycol from urine; the lesions were then chemically reduced back to thymine and thymidine before analysis by HPLC with UV detection (23). The widespread preference for studies of 8-oxodG, however, was derived from 1,000-fold greater sensitivity of 8-oxodG detection using HPLC.
ECD compared with the UV detection of thymine glycol, along with the predominance of 8-oxodG over thymine glycol, in human urine (24). This predominance may arise for reasons, such as the reported instability of thymine glycol and greater levels of 8-oxodG in DNA, due to the relative ease of formation of the latter (24). Overall, this has led to urinary measurements of thymine glycol being superseded by those of 8-oxodG and the subsequent widespread use of HPLC ECD. A number of methods based on column-switching HPLC with regular reverse phase (usually C18) separative columns and ECD for assay of 8-oxodG have been developed (24-28). The pre-purification, or extraction, columns have differed in their packing material, using generally C18 or ion exchange separation approaches. The switching of the effluent from the first to the second column can be controlled automatically by adding to the sample a detectable marker which elutes close 8-oxodG from the first column and the peak is used as the trigger, and this has greatly enhanced the analytic capacity by electrochemical detection (29).

Most recently, mass spectrometric techniques have been increasingly applied to study urinary lesions. Benefits of mass spectrometry include the use of isotopically labeled internal standards, simplifying quantification and accounting for loss during sample workup (and storage), differences in ionization efficiencies due to matrix effects, and confirmation of analyte identity. The HPLC MS/MS assay described by Weimann et al. (21)

### Table 1. DNA markers of oxidative stress reported in the urine of healthy humans

<table>
<thead>
<tr>
<th>Nucleobase</th>
<th>Modification</th>
<th>Abbreviation</th>
<th>Representative method(s) of analysis</th>
<th>Reported levels</th>
</tr>
</thead>
<tbody>
<tr>
<td>Gua</td>
<td>8-oxo-7,8-dihydroguanine</td>
<td>8-oxoGua</td>
<td>HPLC GC/MS (110)</td>
<td>9.4 nmol/mmol Cr</td>
</tr>
<tr>
<td></td>
<td>8-oxo-7,8-dihydro-2'-deoxyguanosine</td>
<td>8-oxodG</td>
<td>HPLC GC/MS (87)</td>
<td>160 nmol/24 h</td>
</tr>
<tr>
<td></td>
<td>8-oxo-7,8-dihydro-2'-deoxyguanosine</td>
<td>8-oxoG</td>
<td>HPLC ECD (110)</td>
<td>~308 pmol/kg/24 h</td>
</tr>
<tr>
<td></td>
<td>Pyrimido[1,2-a]purin-10(3H)-one</td>
<td>M1Gua</td>
<td>LC-MS/MS (21)</td>
<td>48 nmol/24 h</td>
</tr>
<tr>
<td></td>
<td>3-(2-deoxy-D-erythro-pentofuranosyl)-pyrimido[1,2-a]purin-10(3H)-one</td>
<td>M1dG</td>
<td>LC-MS/MS (115)</td>
<td>&lt;LOD (300 pmol/L)</td>
</tr>
<tr>
<td></td>
<td>1,N2-ethenoguanine</td>
<td>1,N2-εGua</td>
<td>LC/APCI-MS/MS (117)</td>
<td>12 fmol/kg/24 h</td>
</tr>
<tr>
<td></td>
<td>N2-3,εethenoguanine</td>
<td>N2-3-εGua</td>
<td>LC/electrospray ionization/MS/MS (60)</td>
<td>&lt;0.3 (LOD)-10 nmol/L</td>
</tr>
<tr>
<td>Ade</td>
<td>8-oxo-7,8-dihydroadenine</td>
<td>8-oxoAde</td>
<td>HPLC GC/MS (33)</td>
<td>7 nmol/L</td>
</tr>
<tr>
<td></td>
<td>8-oxo-7,8-dihydro-2'-deoxyadenosine</td>
<td>8-oxodA</td>
<td>LC-MS/MS (36)</td>
<td>&lt;LOD</td>
</tr>
<tr>
<td></td>
<td>1,N6-etheno-2'-deoxyadenosine</td>
<td>εdA</td>
<td>LC/APCI-MS/MS (56)</td>
<td>38.6 pmol/24 h</td>
</tr>
<tr>
<td></td>
<td>Thymin</td>
<td>Thymidine glycol</td>
<td>HPLC UV (23)</td>
<td>1.9 nmol/mmol Cr</td>
</tr>
<tr>
<td></td>
<td>5-hydroxymethyl</td>
<td>5-HMURa</td>
<td>HPLC GC/MS (79,118)</td>
<td>8.5 nmol/mmol Cr</td>
</tr>
<tr>
<td></td>
<td>5-hydroxymethyl-2'-deoxyuridine</td>
<td>5-HMURa</td>
<td>HPLC GC/MS (118)</td>
<td>At LOD (5 nmol/L)</td>
</tr>
<tr>
<td>Cyt</td>
<td>3,N4-ethenocytosine</td>
<td>εCyt</td>
<td>GC/NICI/MS (119)</td>
<td>84.6 nmol/L/mmol Cr</td>
</tr>
<tr>
<td></td>
<td>3,N4-etheno-2'-deoxycytidine</td>
<td>εdC</td>
<td>3P-postlabeling (120)</td>
<td>2.5 pmol/mmol Cr</td>
</tr>
</tbody>
</table>

Abbreviation: LOD, limit of detection.
was the first report for the simultaneous analysis of the oxidized (8-oxoGua, 8-oxoGuo, and 8-oxodG) and native (Gua, Guo, and dG) moieties. Inclusion of the latter provided useful data for subsequent hypotheses (see Cell Death/ Turnover). Similarly, the HPLC GC/MS assay of Omins’s laboratory entails HPLC prepurification of the compounds of interest, which again includes native compounds namely 8-oxoGua, 8-oxodG, Gua, dG, and 5-(hydroxymethyl)uracil (31, 32), before derivatization and GC-MS. This approach was first applied to the simultaneous analysis of five urinary, oxidatively modified DNA base products, including 8-oxoGua, 5-(hydroxymethyl)uracil, 5-hydroxymethyluracil, 8-oxodG, and 8-oxo-7,8-dihydroadenine (33). Analysis time for HPLC GC/MS is said to be comparable with that of HPLC MS/MS and, at present, is capable of the simultaneous analysis of the greatest number of lesions.

Whereas solid-phase extraction (SPE) had been used to isolate 8-oxodG, a major drawback of many SPE columns is the need to keep them wet during use. The Waters Oasis HLB SPE column, first used by Lin et al. (34), before GC-MS analysis of urinary 8-oxodG, benefits from allowing to become dry during use. These columns have also been used before HPLC MS/MS, which has minimized the incidence of interfering peaks reported by Lin et al. (34) in 10% to 20% of their urine samples (35). The analysis of urinary 8-oxo-7,8-dihydro-2′-deoxyadenosine has been attempted previously (36), but levels were reported to be at or below the limit of detection for the HPLC MS/MS assay (0.3 nmol/L, 7.5 fmol injected, S/N = 3). A similar finding was noted with the SPE-based HPLC MS/MS assay. We predict that SPE will receive increasing application in the future, particularly when on-line SPE is introduced.

**Immunassay.** Competitive ELISA is invariably the format of choice for the analysis of lesions in extracellular matrices, when antibody-based methods are used. Predominantly, 8-oxodG has been the lesion of choice for commercially available [e.g., the Japanese Institute for the Control of Aging (JICA), Stressgen, and Oxis; the latter simply being a distributor of the JICA kit] and custom-made (37, 38) ELISAs. It seems that there are very few commercially available kits presently on the market. Two are available from JICA, named the “new” and “highly sensitive” formats (calibration ranges are 0.5-200 ng/mL and 0.125-10 ng/mL, respectively). These kits are remarkably similar in description (including references cited) to those marketed by Northwest Life Science Specialties, and indeed both utilize the monoclonal antibody denoted N45.1 and have the same concentration range of standards. The other kit available is from Stressgen Bioreagents, now known as Nventa (DNA damage StressXpress ELISA kit), which is remarkably similar to that from Trevigen, Inc. (HT 8-oxodG kit). Both have a range of standards in their calibration curve of 1.9 to 60 ng/mL and essentially the same Microsoft Excel calculation spread-sheet, although the source of the primary antibody used is less clear.

Simply, the assay format of these ELISAs comprise 8-oxodG–protein conjugate bound to the bottom of a 96-well plate, a calibration curve of buffered aqueous 8-oxodG standards containing 0.9% NaCl, and a monoclonal antibody. Both 8-oxodG in the samples and on the plate compete for antibody binding. After a wash step, antibody that remains bound to the plate can be detected using an enzyme-labeled secondary antibody. Color intensity (absorbance) of the enzyme-catalyzed product is proportional to the amount of primary antibody bound, which, in turn, is inversely proportional to the concentration of antigen in the test sample.

The benefits of ELISA are (a) ease of use, (b) no specialist (or indeed expensive) equipment is required, (c) potential application to numerous extracellular matrices (serum, ref. 39; plasma, ref. 40; saliva, refs. 41, 42; urine, ref. 43; cerebrospinal fluid, refs. 44, 45; cell culture medium, refs. 46-48; and sputum, ref. 49), (d) other than centrifugation of cloudy specimens, no pretreatment of urine is required, and (e) high throughput. The kit from JICA has, by far, received the most published use and represents the basis of the bulk of observations to follow. It should be noted that only the JICA kit is recommended for use in plasma; such an application has only recently been reported in the literature, unlike serum.

In the case of plasma, the manufacturer recommends ultracentrifugation of the specimen through a 10-kDa filter (JICA kit insert) to remove “interfering substances,” the identity of which is unknown to us. This has been discussed recently by Cooke et al. (35), who speculated that the competitive format of the ELISA is prone to interference from high molecular weight compounds (e.g., proteins and carbohydrates) present in some body fluids that resulted in high 8-oxodG readings (e.g., serum) or readings in the absence of significant amounts of 8-oxodG (e.g., saliva). Presently, no ELISA kit has been proved to possess sufficient specificity and sufficiently high correlation to the established chemical methods, e.g., HPLC ECD with column switching or HPLC MS/MS.

**Analysis of Secondary DNA Products of Oxidation Reactions in Urine.** Although not as prevalent as methods for analyzing primary DNA products of oxidation in urine, this is a rapidly growing area of interest. Several methods have been developed for the measurement of lipid peroxidation–induced etheno-DNA adducts in human urine. An HPLC fluorescence detection method has been developed for cDA (50) and applied in several studies. For example, in Japanese women, urinary cDA levels were positively associated with increased α-6 polynsaturated fatty acid intake and NaCl excretion, providing evidence of salt-induced inflammation and lipid peroxidation (51). It has also been reported that short-term fasting does not affect urinary cDA adduct or, indeed, 8-oxodG levels in healthy Korean women despite a reduction in primary lipid peroxidation products (e.g., urinary MDA and 8-isoprostaglandin F2α; ref. 52). Isotope dilution GC–negative ion chemical ionization/MS methodology have been developed for 3,N′-etheno-cytosine and 3,N′-ethenodeoxycytidine, with higher levels detected in the urine of smokers compared with...
nonsmokers (53, 54). Furthermore, there was good agreement between urinary levels of 1,N\textsuperscript{2}-ethenoadenine in healthy individuals, measured by HPLC–electrospray ionization–MS/MS method and isotope dilution GC–negative ion chemical ionization/MS methods (55). An HPLC MS/MS method has been developed to analyze urinary 8-oxodG (56) and 3,N\textsuperscript{4}-ethenoxytocytidine (57) and have been applied to show a significant association between etheno-DNA adduct excretion rate and the dietary intake of linoleic acid in healthy men (58).

Background levels of 1,N\textsuperscript{2}-ethenoguanine and N\textsuperscript{2},3-ethenoguanine have been detected in the urine from healthy subjects who have not been exposed to the recognized sources of these adducts, e.g., vinyl chloride and ethyl carbamate (60). An HPLC–electrospray ionization–MS/MS method was developed and used for the detection of low levels of 1,N\textsuperscript{2}-ethenoguanine, which were found to be higher in cigarette smokers (61).

**Method Comparison.** A principal source of criticism of ELISA derives from the discrepancy between chromatography and immunoassay in the determination of "background" or baseline levels of urinary 8-oxodG in healthy individuals. This difference can be anywhere between 4-fold and 10-fold, although recent improvements in the JICA kit (narrowing calibration curve range; recommendation for strict temperature control), have reduced this margin (Fig. 2). For the most part, however, 8-oxodG measurements by the two approaches have shown significant correlation: r = 0.46, P < 0.001 using HPLC ECD (62) and r = 0.73, P = 0.002 using HPLC CC/MS (63); although, this is not always the case (64).\footnote{Cooke et al., unpublished data.}

Nevertheless, this does suggest that both techniques share a common analyte, with the ELISA perhaps recognizing additional compounds. In accounting for the discrepancy, some reports in the literature fail to distinguish between antibodies and their respective specificities, applying a blanket statement to the unsuitability of all ELISAs. Whereas some of this criticism is well-founded, clearly some antibodies are highly specific and their application to urine seems entirely appropriate.

For example, according to characterization data reported by Toyokuni et al. (65), N45.1 is highly specific for 8-oxodG. In addition to being specific for the hydroxylated C8 position of Gua, it would seem to discriminate the C6 carbonyl group of 8-oxoGua (and C2 NH\textsubscript{2}) from the C6 amino group of 8-oxo-7,8-dihydroxyadenine. Furthermore, unlike many antibodies, N45.1 displays preference for the 2-deoxyribose moiety of 8-oxodG until recently, a prerequisite to the assessment of urinary DNA oxidation products. The closest competitor for the antibody is 8-oxo-7,8-dihydroguanosine (8-oxoG), which needs to be present in concentrations two orders of magnitude higher than 8-oxodG to compete to the same extent. Weimann et al. (20) later showed that 8-oxoG was present at a concentration not dissimilar to 8-oxodG itself (8-oxoG, 48 nmol/24 h versus 8-oxoG, 28 nmol/24 h), and hence, it was concluded that 8-oxoG was unlikely to be a competitor (66).

Similarly, whereas 8-oxoGua is present in urine at significant concentrations (136 ± 12 nmol/24 h), characterization data suggests reactivity of N45.1 toward the modified base to be, at worst, negligible. Some of the comments in the literature, which state that N45.1 recognizes 8-oxoGua, are likely derived from misinterpretation of a statement in the abstract of the paper by Toyokuni et al. (65), "N45.1 recognized both the modified base and 2-deoxyribose structure of 8-oxodG." Clearly, some authors have taken this to refer to the modified base and 2-deoxyribonucleoside as separate structures, not in combination, as intended by Toyokuni et al. (65). It is important to note that whereas the JICA assay is reported to be specific for 8-oxodG, the specificity of the Trevigen/Stressgen kit(s) includes recognition of 8-oxoGua and 8-oxoG, likely to account for the higher levels of urinary 8-oxodG reported by these kits. The appropriateness of a calibration curve consisting solely of 8-oxodG, as used by these latter two suppliers’ kits, is also questionable.

Identifying the basis of the discrepancy has proved to be problematic. We have speculated that, as chromatographic assays can only detect "free" 8-oxodG, other compounds structurally related to 8-oxodG, recognizable by N45.1 specifically, may exist and be present in urine (67). For example, 8-oxodGMP, a product of the activity of enzymes, such as the human Mut T homologue (hMTH1; ref. 68), which may have the potential to leave the cell (69); along with 8-oxodG-containing oligomers, putative products of nucleotide excision repair (67). The issue 8-oxodG containing oligomers in urine does not seem to have been resolved, as there have been two reports both indicating (70) and disputing their presence (71), although both reports agree that oligomers are present. Patel et al. (70) ascribe the failure of Weimann et al. (71) to detect 8-oxodG–containing oligomers to a sensitivity issue, arising from the absence of sample concentration steps despite LC-MS/MS having greater than thrice the sensitivity of the ELISA. At best, this would indicate that, if present, 8-oxodG–containing oligomers are not at a concentration detectable in urine by the ELISA and hence not the source of the discrepancy between this and chromatographic techniques. Furthermore, use of an independent technique, i.e., not one...
relying upon antibody specificity, would provide a more definitive answer.

Reardon et al. (72) showed that 8-oxodG is a substrate for nucleotide excision repair, which excises lesions as a single-stranded, lesion-containing oligomer (see DNA Repair). Galloway et al. (73) showed that, postexcision, cyclobutadithymine (T<->T)-containing oligomers, at least, are subsequently degraded to lesion-containing 7-mers. What then happens to the oligomers is unknown, although from the findings of Le Curieux and Hemminki (74), who detected the presence of urinary T<->T as a monophosphate nucleotide dimer, it may be implied that further processing does occur, the nature of which remains to be described. Despite all this circumstantial evidence, the precise reason for the higher ELISA values remains unknown and represents a severe problem that needs to be clarified before its use can be recommended without comparison with established chromatographic methods.

The HPLC ECD method is not without analytic difficulties, Bogdanov et al. (30) reported the presence of peaks which coelute with 8-oxodG identified by multichannel colometric electrochemical detection. Furthermore, if DNA oxidation products are indeed present in urine in forms other than bases or 2’-deoxyribonucleosides, such as oligomers, monophosphates, or diphosphates, then the vast majority of chromatographic methods will not recognize these products. Despite these issues, within technique agreement of levels in control subjects between laboratories is strong and supports the continued use of both chromatographic and ELISA approaches. Furthermore, in longitudinal or comparative studies, the significance of absolute levels is of less importance compared with the ability to detect differences in 8-oxodG levels.

**Urinary Lesions: Issues of Artifact and Stability.** The measurement of oxidatively generated damage to DNA has been plagued by the issue of artifact-generated, for example, during DNA extraction, or sample workup. This issue has been critically examined in detail, European Standards Committee on Oxidative DNA Damage, as part of its attempts to achieve consensus between different techniques and different laboratories (the objectives of European Standards Committee on Oxidative DNA Damage are outlined in Table 2). Similarly, urinary lesions may arise from the artifactual oxidation of DNA or its constituent nucleobases or 2’-deoxyribonucleosides after exposure to metabolic enzymes or other oxidizing species after release into the systemic circulation or in the urine. However, Shigenaga et al. (75) showed that dG is not artifactual oxidized in the systemic circulation or after incubation with microsomal enzymes. Equally, 8-oxodG, thymine glycol, and thymidine glycol are not subject to degradation upon release into the systemic circulation (23, 75), the stability of 8-oxoGuA not having been examined. Experimental studies with 8-oxodG injected i.v. have shown that recovery is almost complete in urine in pigs, indicating that further metabolism is an unlikely event (76).

Furthermore, as significant concentrations of hydrogen peroxide have been reported to be present in urine (77), Lin et al. (34) examined the potential for 8-oxodG to be generated from dG present in the urine. Combinations of dG (1 nmol) and hydrogen peroxide (50 or 100 μmol/L) were incubated in a number of urine samples for 24 h at room temperature (34). The authors concluded that hydrogen peroxide has a negligible effect upon 8-oxodG production, under these circumstances, agreeing with previous results from Ames’ laboratory.

As part of a study, representing a great deal of foresight, Lotf et al. (78) reported that storage of urine samples at ~20°C and repeat measurements did not lead to a decrease in the concentration of 8-oxodG over a 15-year period, confirming the stability of this lesion in urine. This also applies to 5-(hydroxymethyl)uracil that has been stable upon storage at ~80°C for periods up to 4 months (79).

Whereas the above provide evidence of stability in extracellular matrices, it is worth noting that 8-oxodG is more prone to oxidation than dG due to its lower redox potential (80, 81) and is preferentially oxidized even in the presence of an excess of Gua (82, 83). By the same token, oxidation of Thy to thymine glycol is less likely to occur than Gua to 8-oxoGua due to the lower oxidation potential of Gua compared with Thy. However, it would seem that most of these studies have examined the oxidation of 8-oxoGua *in situ* in DNA (84), rather than as a postexcision product of DNA repair, which may alter the likelihood of oxidation and its “oxidizability.” In contrast, M1G and 3-(2-deoxy-β-D-erythro-pentofuranosyl)pyrimido[1,2-a]purin-10(3H)-one both seem to undergo further oxidative metabolism in rat liver cytosol, with the base adduct being a better substrate for such enzymic oxidation than the deoxyribonucleoside adduct (85, 86). There is also some evidence to suggest that M1G is further oxidized when given i.v., although 3-(2-deoxy-β-D-erythro-pentofuranosyl)pyrimido[1,2-a]purin-10(3H)-one was not examined (85).

Overall, it would seem that 8-oxodG, thymine glycol, and thymidine glycol, and probably 8-oxoGuA, are not formed artifically *in vivo* in biological matrices of mammals, including humans, and there is no published evidence for their degradation upon release. In contrast, there is strong evidence that M1G and 3-(2-deoxy-β-D-erythro-pentofuranosyl)pyrimido[1,2-a]purin-10(3H)-one may undergo oxidation, either enzymically or in the systemic circulation.

**Sources of Extracellular, Oxidatively Modified DNA Lesions**

Broadly speaking, the possible sources of extracellular, oxidatively modified DNA lesions are (a) diet, (b) cell death/tturnover (mitochondrial turnover), and (c) DNA repair.

**Diet.** Urine is the extracellular matrix upon which dietary influence has been most studied. The majority of reports measuring oxidatively modified DNA lesions in urine have focused upon 2’-deoxyribonucleoside lesions and 8-oxodG specifically. This is largely a consequence of early work from Bruce Ames’ laboratory in rats, showing that diet could affect levels of urinary 8-oxoGuA, but not 8-oxodG, and they concluded thymine glycol to be unaffected by diet (discussed in detail in ref. 67). There is little human data on the subject of dietary contribution and some disagreement with the animal findings. Gackowski et al. (87) have presented evidence to suggest that, in humans, neither 8-oxoGuA nor 8-oxodG is
The presence of [15N5]-8-oxodG and [15N5]-8-oxoGua in urine was examined using HPLC and radiolabeled lesions are preferred over those involving nucleic acid–free diets as the use of
radiolabels provides a degree of stringency not afforded by nucleic acid–free diets. However, for ethical reasons, radiolabeled lesions are to be avoided in human studies. To circumvent this issue, Cooke et al. (32) recently completed a feeding study using heavily oxidized, stable isotopically labeled [15N]-DNA (98% incorporation). Healthy male volunteers were fed oxidized [15N]-DNA, and first void, midstream urine samples were collected for up to 14 days later. The presence of [15N5]-8-oxodG and [15N5]-8-oxoGua in urine was examined using HPLC. Neither lesion was detected in any of the urine samples. To confirm that the absence of lesion was not a sensitivity issue, urinary [15N5]-8-oxoG and [15N5]-8-oxoGua, which should be present at significant levels, was also examined. Again, neither of these labeled compounds were detected in any of the urine samples. The authors also investigated whether any of the ingested [15N5]-Gua/[15N5]-dG had been degraded to [15N5]-uric acid, the final product of purine metabolism, before excretion.

Again, no labeled material was detected. As with previous studies (75), the authors could only presume that the isotopically labeled DNA components/adducts pass through the gastrointestinal tract to appear in feces. Nevertheless, these results, coupled with the findings of Gackowski et al. (87), provide the most compelling evidence that dietary [15N5]-DNA is absorbed and excreted via the urine in vivo.

To achieve consensus between above methods
Investigate any discrepancy between methods
Provide validated methods for wider dissemination
Establish a reference range for urinary 8-oxodG, accounting for factors known to affect levels
Investigate the use of other oxidized nucleobases and nucleosides as possible biomarkers
Elucidate extent of contribution from possible confounders of urinary lesion measurement (e.g., diet and cell turnover)

Cell Death/Turnover. In 1993, it was asserted that urinary 8-oxodG did not reflect DNA repair, as it was not a product of base excision repair (90) rather it was a product of non-specific nucleases acting upon DNA released from during cell death, liberating dG, which is subsequently oxidized. This discounted the potential existence of other, as yet undiscovered, repair pathways which may yield 8-oxodG. Whereas repair processes, whose reaction product may be 8-oxodG, have been described (see below), the possible contribution from cell turnover remains to be addressed, although some data are beginning to emerge. For example, it has been reported recently that rat liver homogenates can release 8-oxodG from oxidatively modified DNA (91). The authors suggest that, during DNA degradation, most of the 8-oxodG, present in DNA, is released by enzymes and subsequently appears in the urine, implying this to be a major source. However, they do acknowledge that their system does not accurately represent the in vivo situation, not in the least as it is unrealistic — multiple uncompartmentalized nucleases acting upon circulating DNA, particularly as the majority of cell death in healthy humans will be via apoptosis.
The evidence against a contribution from cell turnover has, for the most part, been anecdotal. There exists a number of reports in which urinary 8-oxodG has been measured in patients undergoing chemotherapy in the absence of concomitant increase in urinary 8-oxodG (reviewed in ref. 66). As yet, the most decisive argument against the contribution of cell death to urinary levels of 8-oxodG and 8-oxoGua comes from a recently published report from the Olinski group (92). The findings from this study are supported by the results of Weimann et al. in a study examining oligonucleotides in urine, which concluded that the limited excretion of oligonucleotides into urine argues against oligonucleotides or, indeed, nucleosides originating from cell death (71). On balance, these results suggest that the contribution of cell death to urinary 8-oxodG levels is minimal, but clearly further work needs to be done.

**DNA Repair.** With respect to DNA repair, the source of oxidatively modified DNA bases in extracellular matrices seems clear. The DNA N-glycosylases responsible for removing modified bases are increasingly well defined (93) and have been discussed thoroughly elsewhere (2). 8-Oxoguanine DNA glycosylase 1 (OGG1) is the major enzyme involved in the removal of 8-oxoGua from cellular DNA. Recently, levels of 8-oxoGua were determined in OGG1−/− mice and compared with a wild-type strain. There was ~26% reduction in levels of urinary 8-oxoGua in the deficient strain compared with the wild type, and no significant changes in 8-oxodG were observed (94). This indicates that mouse OGG1 glycosylase is a significant, but by no means unique, source of urinary 8-oxoGua.

The results clearly suggest the existence of back-up DNA glycosylase(s) that cannot entirely compensate for OGG1 deficiency. The reversion to a less efficient back-up system should result not only in reduction of 8-oxoGua in urine but is also expected to lead to the simultaneous, gradual increase in the background level of 8-oxoGua in cellular DNA (discussed fully in ref. 95). In female OGG1−/− mice of 8 to 10 weeks of age, no significant increase in 8-oxoGua, assessed as formamidopyrimidine DNA glycosylase–sensitive sites, was found in the liver compared with wild-type mice, whereas the level in lung tissue was 3-fold greater than wild type (96). The faster accumulation in lung could be due to high level of oxygen exposure compared with liver.

The OGG1−/− mice also accumulate much more 8-oxodG in target organs after exposure to oxidative stress-inducing agents, such as diesel emission particles or potassium bromate. In the mitochondrial genome from liver tissue, a 20-fold increase in the 8-oxoGua level was observed in the OGG1 defective mice compared with wild type (97). Taken together, these data suggest the existence of another repair pathway, the product of which is not 8-oxoGua or indeed 8-oxodG that maintains lowered 8-oxoGua levels in most tissues even in the absence of OGG1. It is worth considering that the rate of 8-oxoGua accumulation in DNA may be determined by the proliferation rate of the tissue so that rapid proliferation may dilute the effect of OGG1 deficiency, whereas 8-oxodG accumulates in liver, lung, and muscles with a low proliferation rate.

The presence of 2′-deoxyribonucleoside lesions in extracellular matrices is very much less well defined, as there are no reports of a single-DNA repair enzyme whose activity yields 8-oxodG. Based upon existing evidence, we have proposed the following three DNA repair routes (Fig. 3) as the most likely contributors to the presence of oxidatively modified 2′-deoxyribonucleosides in urine.

**Nudix Hydrolases.** There is an imperative for preventing modified DNA precursors from being incorporated into the genome. The best characterized enzyme which performs such a role is the 8-hydroxy-2′-deoxyguanosine triphosphatase (68) activity of NUDT1 (MutT homologue, MTH1), hydrolyzing 8-hydroxy-2′-deoxyguanosine triphosphatase to 8-oxodGMP (Fig. 3). It has been suggested that further processing, perhaps by 5′(S′)-nucleotidases, may give rise to 8-oxodG, which is not charged and can be removed from the cell, ultimately appearing in the urine (68). Recent data from the group of Harms-Ringdahl strongly imply that NUDT1 activity and, thus, the nucleotide pool represent a major source of extracellular 8-oxodG (98). These studies in cell culture used small interfering RNA gene expression knockdown methodology directed at human NUDT1, however, the roles of other Nudix hydrolases, such as NUDT15 (MTH2) and NUDT5, which include 8-hydroxy-2′-deoxyguanosine triphosphatase and 8-oxodGDP among their substrate repertoire, respectively, remain to be defined (99, 100). These new observations on the contribution of NUDT1 to extracellular 8-oxodG, coupled with any potential negation of the role of diet and cell turnover, could mean that we are very close to determining the biological significance of urinary 8-oxodG levels as markers of oxidation in and sanitization of the nucleotide pool.

**Nucleotide Excision Repair.** Despite seeming to be directed principally toward bulky lesions, such as cyclobutane thymine dimers (T<>T), there is some evidence that nucleotide excision repair (Fig. 3) may act upon nonbulky lesions, such as 8-oxoGua (72). Indeed, the rate of 8-oxoGua removal seems comparable with that for T<>T (72), seemingly generating a lesion-containing oligomer, ~24 to 32 nucleotides long (101). A recent report failed to show 8-oxoGua−containing oligomers in urine (71), implying that further processing occurs, perhaps ultimately yielding 8-oxodG, or that they do not exist. However, under normal circumstances, the role of nucleotide excision repair in the removal of 8-oxoGua and perhaps other small oxidatively generated DNA lesions would seem to be negligible (102-106), although results from xeroderma pigmentosum cell lines have not entirely excluded the possibility (72, 103, 107, 108).

**Endonuclease(s).** A poorly characterized endonuclease has been reported by Bessho et al. (109), which, lacking a glycosylase activity, is predicted to give rise to 3′,5′-8-oxodGDP as the putative product. We have previously proposed that this may be subsequently hydrolyzed to 8-oxodG by nucleotidase(s). The recently described nucleotide incision repair process generates lesion-containing X’dNMP products; again, these could be subsequently metabolized to a 2′-deoxyribonucleoside. However the activity of nucleotide incision repair toward 8-oxodG−containing substrates is reported to be negligible. Whereas this may be the case, it is entirely feasible that this process could be a potential source of other extracellular lesion-containing 2′-deoxyribonucleosides.

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Although the types of lesion that have been examined as substrates have yet to be detected in urine.

**European Standards Committee on Urinary (DNA) Lesion Analysis**

There is a pressing need for (noninvasive) biomarkers of oxidative stress that are sufficiently validated to be applied to large-scale, molecular epidemiology studies. Validation should include a detailed understanding of the biomarker's provenance plus robust and reproducible methods for analysis. The noninvasive nature of such biomarkers minimizes ethical issues and allows studies to incorporate vulnerable subjects, such as the very young and the very old. Furthermore, the biomarkers should have long-term stability under routinely used conditions, reducing the need for new, prospective studies by allowing retrospective analysis of existing cohorts. Widespread agreement between methods for biomarker analysis will allow multiple accredited centers to perform analyses, the results of which may hence be pooled and reduce the burden on any one laboratory.

Urinary 8-oxodG is an ideal candidate for fulfilling these criteria. European Standards Committee on Urinary (DNA) Lesion Analysis (ESCULA) has been formed to complete the validation of this biomarker of oxidative stress. As this review indicates, the origin of 8-oxodG in urine is perhaps not as simple as first thought. ESCULA is currently addressing this issue through $[15N]_{2}$-labeled DNA feeding studies, experiments incorporating knockout mice and markers of cell turnover. As noted above, there is a discrepancy in basal urinary 8-oxodG levels when comparing chromatographic techniques with ELISA, although all techniques have been shown to discriminate between diseased and healthy subjects. ELISA has received widespread use and is clearly amenable to the greatest number of laboratories; however, this discrepancy continues to raise questions regarding its utility. To continue the validation of urinary 8-oxodG, ESCULA is presently conducting a multicenter study, incorporating numerous platforms and methodologies, including ELISA (see Table 2). We propose that understanding the basis of this discrepancy will aid our understanding of the significance of urinary lesions. An additional outcome of this interlaboratory study will be the provision of validated assays for urinary 8-oxodG measurement for widespread dissemination and application. Hitherto, this has been done in a limited fashion, and the discrepancies remain unaddressed. ESCULA will also provide advice concerning best practice, for example, serious consideration needs to be given to how results for urinary lesion measurements are expressed and the applicability of correcting for creatinine. Finally, there is growing clinical interest in the measurement of urinary 8-oxodG as a means to determine the role of oxidative stress in disease and evaluate intervention strategies. With assays validated, numerous laboratories may work together to establish reference ranges for normal and pathologic conditions and conduct the new molecular epidemiologic studies.

**Acknowledgments**

ESCULA wishes to dedicate this review to the memory of the late Dr. Jagadeesan Nair (DKFZ, Germany), who passed away in late 2007.

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**Figure 3.** DNA repair sources of extracellular 8-oxoG and 8-oxoGua. NER, nucleotide excision repair; hOgg1, human 8-oxoguanine glycosylase.
12 Urinary Oxidatively Modified DNA Lesions


Urinary Oxidatively Modified DNA Lesions

Cai JP, Ishibashi T, Takagi Y, Hayakawa H, Sekiguchi M. Mouse

Haghoost S, Sjolander L, Czen S, Harms-Ringdahl M. The

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Measurement and Meaning of Oxidatively Modified DNA Lesions in Urine

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