DNA strand breaks and base modifications induced by cholesterol hydroperoxides

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Abstract

Cholesterol (Ch) can be oxidized by reactive oxygen species, forming oxidized products such as Ch hydroperoxides (ChOOH). These hydroperoxides can disseminate the peroxidative stress to other cell compartments. In this work, the ability of ChOOH to induce strand breaks and/or base modifications in a plasmid DNA model was evaluated. In addition, HPLC/MS/MS analyses were performed to investigate the formation of 8-oxo-7,8-dihydro-2'-deoxyguanosine (8-oxod-Guo) after the incubation of 2'-deoxyguanosine (dGuo) with ChOOH and Cu²⁺. In the presence of copper ions, ChOOH induced DNA strand breaks in time and concentration-dependent manners. Purine and pyrimidine base modifications were also observed, as assessed respectively by the treatment with Fpg and Endo III repair enzymes. The detection of 8-oxodGuo by HPLC/MS/MS is in agreement with the dGuo oxidation in plasmid DNA. ChOOH-derived DNA damage adds further support to the role of lipid peroxidation in inducing DNA modifications and mutation.

Keywords: Cholesterol hydroperoxides, DNA damage, 8-oxo-7,8-dihydro-2'-deoxyguanosine, HPLC/MS/MS.

Introduction

Cholesterol (Ch) is an essential lipid constituent of biological membranes. However, it can be oxidized under a variety of conditions, forming oxidized products such as Ch hydroperoxides (ChOOH). Lipid hydroperoxides (LOOH) are prominent intermediates of peroxidative reactions induced by activated oxygen species such as hydroxyl radical, lipid alkoxyl or peroxyl radicals, singlet molecular oxygen ($^{1}O_{2}$) and peroxynitrite [1,2]. Once formed, they can be decomposed by heating, exposure to UV light (in some cases) or by the addition of transition metals [3]. Decomposition of peroxides can generate alkoxyl and peroxyl radicals that may be able to oxidize other molecules such as proteins and DNA.

The effects of DNA adducts generated by aldehydes derived from lipid peroxidation have been extensively studied [4–7]. In addition, many efforts have been made to evaluate the role of LOOH in the induction of

DNA damage, since they are implicated in the aetiology of human cancers and mutations [2,8]. For example, it has been shown that linoleic [9-11] and arachidonic acid [12,13] hydroperoxides are able to induce strand breaks and base modifications. In this sense, ChOOH are of special interest, since they were shown to be able to translocate among LDL and membranes of cell, nuclei and organelles faster than the parent Ch [14]. Moreover, unlike other LOOH, ChOOH are completely resistant to classical GSH peroxidase (GPx-1) catalysed reduction. Phospholipid hydroperoxide GSH peroxidase (GPx-4) is the only known enzyme able to detoxify ChOOH, although more slowly than other hydroperoxides [15]. These characteristics make ChOOH potentially more dangerous since they may be able to disseminate peroxidative stress.

Despite the potential role of ChOOH in propagating peroxidative stress through different cell

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compartments, the ability of these hydroperoxides to induce DNA damage has not been evaluated so far. In addressing this issue, we have examined the ability of ChOOH to induce strand breaks and base modifications in plasmid DNA. Moreover, 8-oxo-7,8-dihydro-2'-deoxyguanosine (8-oxodGuo) was detected by HPLC/MS/MS analyses, in agreement with 2'-deoxyguanosine (dGuo) oxidation in plasmid DNA.

Materials and methods

Materials

Cholesterol, dGuo and all the other chemicals were from Sigma (St. Louis, MO) and the solvents were from Merck (Darmstadt, Germany). Formamidopyrimidine-DNA glycosylase (Fpg) and Endonuclease III (Endo III) were purchased from New England BioLabs (Beverly, MA). All the solutions were prepared with distilled water purified with a Millipore Milli-Q system (Bedford, MA).

Cholesterol hydroperoxides synthesis, purification and quantification

ChOOH were synthesized by photo-oxidation using methylene blue as a sensitizer. Typically, 100 mg of Ch were dissolved in 50 mL of chloroform containing 0.2 mM of methylene blue and irradiated with a tungsten lamp (500 W) for 3 h. Irradiation was conducted in an ice-bath under a continuous flux of oxygen. The methylene blue was removed and ChOOH were isolated by silica gel column chromatography. Briefly, the mixture was placed in the column and isolated according to the following *n*-hexane (H) diethyl ether (E) gradient: 10 mL H; 20 mL H:E (80:20); 50 mL H:E (70:30); 50 mL H:E (65:35); 30 mL H:E (60:40); 30 mL H:E (55:45); 30 mL H:E (50:50). The concentration of ChOOH was determined by iodometry [16].

Cholesterol hydroperoxides analysis by HPLC/MS

Cholesterol hydroperoxides were analysed by HPLC (Shimadzu, Kyoto, Japan) coupled to a Quattro II Micromass mass spectrometer (Manchester, UK), with an atmospheric pressure chemical ionization (APCI) source in the positive ion mode. The ChOOH were eluted using a 250 mm \times 4.6 mm (particle size 5 µm) Luna C-18 reverse-phase column (Phenomenex, Torrance, CA, USA) using 95% methanol and 5% water at a flow rate of 1 mL/min. The UV detector was set at 210 nm. For MS analyses, a small fraction of the eluant was directed into the mass spectrometer at a flow rate of 100 µL/min. The source and desolvation temperatures were held

at 150°C and 300°C, respectively. The flow rates of drying and nebulizing gas were optimized at 300 and 15 L/h, respectively. The cone voltage was set at 20 V and the corona potential at 3.4 kV. Full scan data were acquired over a mass range of 100–700 m/z.

Cholesterol hydroperoxides analysis by NMR

NMR analyses were recorded in DPX300 instrument (Bruker-Biopsin, Rheinstetten, Germany). ChOOH were identified as 3β -hydroxycholest-5-ene-7 α hydroperoxide (7α-OOH), 3β-5α-cholest-6-ene-5hydroperoxide (5 α -OOH), 3 β -hydroxycholest-4ene-6a-hydroperoxide (6a-OOH) and 3β-hydroxycholest-4-ene-6\u00df-hydroperoxide (6\u00bf-OOH) comparing chemical shifts with previous works [17-19]. 7 α -OOH – ¹H NMR (300 MHz, CDCl₂) δ ppm 5.72 $(dd, \mathcal{J} = 4.94, 1.74 Hz, 1H), 4.16 (dt, \mathcal{J} = 4.64)$ 4.60, 1.72 Hz, 1H), 3.68–3.56 (m, 1H). 5α-OOH – ¹H NMR (300 MHz, C_6D_6) δ ppm 5.70 (dd, $\mathcal{J} =$ 10.01, 0.91 Hz, 1H), 5.55 (dd, f = 9.97, 2.03 Hz, 1H), 4.19–4.02 (m, 1H). 6α -OOH – ¹H NMR (300 MHz, CDCl₃) δ ppm 5.59 (q, $\mathcal{J} = 1.69, 1.69,$ 1.69 Hz, 1H), 4.49 (tdd, $\mathcal{J} = 12.28$, 4.63, 1.79, 1.79 Hz, 1H), 4.26–4.13 (m, 1H). 6β -OOH – ¹H NMR (300 MHz, CDCl₃) δ ppm 5.66 (t, f = 1.69, 1.69 Hz, 1H), 4.33 (dd, $\mathcal{J} = 3.99$, 2.26 Hz, 1H), 4.25-4.13 (m, 1H).

Incubation of plasmid pBSK-II DNA with ChOOH in the presence of Cu^{2+} followed by incubation with DNA glycosylases

Supercoiled plasmid pBSK-II DNA was prepared according to standard methods and purified using QIAGEN[®] kit (Valencia, CA). DNA reactions with ChOOH consist of 1.4 µg of DNA, 0.5-10 mM of ChOOH and 10 µM of Cu2+ in 30 mM of sodium phosphate buffer pH 7.4 (final volume 100 µL). Control experiments were run replacing ChOOH solution by ethanol. For solubility reasons, all reactions contain 30% ethanol. Reaction mixtures were incubated for selected times at 37°C under agitation (conditions are depicted in figure legends). After that, ChOOH were extracted from the reaction mixtures with 300 µl of diethyl ether (repeated twice). Subsequently, 10 μ L of the reaction containing the DNA were incubated with 0.8 U of Fpg (10 mM Bis Tris Propane-HCl (pH 7.0), 10 mM MgCl₂, 1 mM dithiothreitol and 0.1 mg/mL bovine serum albumin) or 1 U of Endo III (20 mM Tris-HCl (pH 8.0), 1 mM EDTA, 1 mM dithiothreitol and 0.1 mg/ mL bovine serum albumin). Control experiments were done replacing enzyme solution by water. Incubations were carried out at 37°C for 30 min. After that, DNA was mixed with 2 µL of loading buffer (0.25% bromophenol blue and 15% Ficoll

type 400) and loaded onto a 1% agarose gel containing 0.3 μ g/mL ethidium bromide. The gels were run at a constant voltage of 100 mV for 120 min in 90 mM Tris/borate 2 mM EDTA buffer (pH 8.0). The gels were photographed and quantified using an Epi Chemi II Darkroom (Bio Imaging Systems, Upland, CA, USA) and a Lab WorksTM Image Acquisition and Analysis Software (UVP, Upland, CA). Results represent the average and SD of, at least, three independent determinations.

Incubation of 2'-deoxyguanosine with ChOOH

Reactions of dGuo with ChOOH consisted of 1 mM dGuo, 10 mM ChOOH and 100 μ M Cu²⁺ (final volume 100 μ L). Control experiments were run replacing ChOOH solution by ethanol. In order to maximize the interactions between dGuo and ChOOH, the incubations contained 70% of ethanol. The samples were incubated for selected times at 37°C under agitation. After that, 7 pmol of [M+5] isotopically labelled 8-oxodGuo were added to the samples as an internal standard. Then, ChOOH were extracted from the reactions with 300 μ l of diethyl ether (repeated three times). Subsequently, extracted samples were dried under a N₂ atmosphere, re-suspended in water and injected into the HPLC/MS/MS system.

Determination of 8-oxo-7,8-dihydro-2'-deoxyguanosine by HPLC/MS/MS

Samples resulting from dGuo reactions with ChOOH were analysed by HPLC/MS/MS (Shimadzu HPLC coupled to Quattro II Micromass mass spectrometer). The reaction products were separated in an LC-18 column (Phenomenex, 150×2.1 mm, 5 µm) using an isocratic gradient of ammonium formate (10 mM, pH 3.7): acetonitrile (96:4) at a flow rate of 0.1 mL/ min. The eluent was monitored by electrospray

ionization (ESI-MS) in the positive ion mode. The source and desolvation temperatures were held at 100°C and 150°C, respectively. The flow rates of drying and nebulizing gas were optimized at 300 and 15 L/h, respectively. The cone voltage was set at 15 V and the collision energy was set at 15 eV. Full scan data were acquired over a mass range of 100–500 m/z. The quantification was performed in the selected reaction monitoring mode (SRM), by selecting the mass transition of the 284 to 168 for 8-oxodGuo and 289 to 173 for the internal standard ([M+5] isotopically labelled 8-oxodGuo). Isotopically labelled 8-oxodGuo was a gift of Dr. Jean-Luc Ravanat.

Results

Cholesterol hydroperoxides analysis by HPLC/MS

After synthesis, ChOOH were purified from the parent Ch and other oxidized products, with purity and identities checked by HPLC/MS and NMR analyses. Figure 1 shows four product peaks identified in order of elution as 7α -OOH, 5α -OOH, 6α -OOH and 6β -OOH. Chloroform was used as the solvent in the photosensitization. This fact accounts for the lower yield of 5 α -OOH compared to 7 α -OOH, since it is well known that 5α -OOH re-arranges to 7α -OOH in non-polar solvents [17,20]. A MS spectrum from 5α-OOH, representative for all isomers, is displayed in Figure 1C. Protonated molecule (m/z 419) is not seen in the MS spectrum. However, the detection of fragments derived from protonated molecules indicates that the loss of the first H₂O molecule is a favoured reaction. In fact, the characteristic fragmentation pattern of ChOOH is the consecutive loss of H₂O molecules, yielding cationic fragments at m/z 401, 383 and 367. These results are in agreement with previous analyses for ChOOH [21,22]. A mixture of the four ChOOH isomers was used to test the possibility of DNA damage generation.



Figure 1. HPLC/APCI/MS analyses of ChOOH. (A) UV detection at 210 nm. (B) MS chromatogram selection m/z 401. (C) Mass spectrum at 15.44 min. Numbers over the peaks in (A) and (B) correspond to the retention times (min) in order of elution for 7α -OOH, 5α -OOH, 6α -OOH and 6β -OOH.

Induction of strand breaks was assessed by agarose gel electrophoresis. In each lane, the upper band corresponds to the open-circular DNA (OC), the intermediate band (if present) to linear DNA (L) and the lower band to supercoiled DNA (SC). For each lane, the SC and OC forms were summed up and the extent of DNA oxidative damage was expressed as the percentage of DNA SC form. A correction factor of 1.4 was applied to the value of SC forms to account for their fluorescence quantum yield lower than the OC form [23]. All reaction mixtures contained 30% ethanol, to allow ChOOH solubilization. The exposure of plasmid DNA to increasing concentrations (0.5-10 mM) of ChOOH in the presence of 10 μ M of Cu²⁺ led to an increase in DNA damage, as shown by the decrease in the SC DNA form (Figure 2A₁). A plot of these results is shown in Figure 2B. For all experiments, SC amount found in control incubation was considered as 100% and the results were reported as a control percentage. In this way, after 1 h of incubation with 0.5 mM of ChOOH, the SC DNA percentage had only a slight decrease to 97%. However, increasing ChOOH concentration enhances the strand breaks level, with the SC DNA percentage decreasing to 82% upon exposure to 10 mM of ChOOH. Control experiments using Cu²⁺ alone (Figure 2A₁, lane 1) or ChOOH in the absence of Cu²⁺ (data not shown) have shown no significant strand scission.

Besides strand breaks, ChOOH-decomposition products could also lead to base oxidation, resulting in products which are potential substrates for base excision repair (BER) enzymes. In order to test this possibility, we use two BER enzymes, Fpg and Endo III. These proteins possess both DNA glycosylase and apurinic/apyrimidinic (AP) lyase activities, creating additional breaks at sites of oxidized bases. Some of the damaged bases recognized by Fpg are 8-oxodGuo, 2,6-diamino-4-hydroxy-formaminopyrimidine and a broad spectrum of modified purines [24,25]. On the other hand, Endo III recognizes and removes a number of thymine- and cytosine-derived lesions in DNA [26,27].

Treatment of plasmid DNA with Fpg after 1 h of ChOOH exposure led to a significant decrease in SC DNA form, consistent with purine bases oxidation (Figure 2A₂). Upon exposure to 0.5 mM of ChOOH, SC DNA percentage decreased only to 90%, although upon exposure to 10 mM of ChOOH the same percentage decreased to 71% (Figure 2B). Interestingly, even untreated DNA contains a small amount of modified bases (data not shown). This observation is in agreement with a previous work [28]. Neither the treatment of plasmid DNA with Cu^{2+} alone (Figure 2A₂, lane 1) nor the treatment with ChOOH in the absence of Cu^{2+} (data not shown) were able to significantly increase DNA damage as compared with untreated DNA. Contrasting with Fpg treatment, incubation of plasmid DNA with Endo III leads just to a slight increase in the pre-existing damage on the ribose phosphate



Figure 2. Induction of strand breaks upon exposure of pBSK-II plasmid DNA to ChOOH (0.5–10 mM) and 10 μ M of Cu²⁺. Reactions mixtures were incubated for 1 h at 37°C in phosphate buffer (pH 7.4, with 30% ethanol, to allow ChOOH solubilization). Subsequently, ChOOH were extracted and DNA was subjected to a second incubation with DNA glycosylases for 30 min at 37°C. (A) Agarose gel electrophoresis. (1) Incubation in the absence of enzymes; (2) Incubation in the presence of 0.8 U of Fpg; (3) Incubation in the presence of 1 U of Endo III. Lane 1: control reaction in the absence of ChOOH; Lanes 2–7: incubation with 0.5, 1.0, 2.5, 5.0, 7.5 and 10 mM of ChOOH, respectively. Plasmid DNA forms: OC: open circular; SC: supercoiled. (B) Control percentage of supercoiled DNA. The SC percentage was calculated considering the SC amount of control incubation as 100%. Results represent the average and SD of at least three independent determinations.

backbone (Figures $2A_3$ and B), showing that pyrimidine bases are less susceptible than purines to ChOOH-derived oxidation.

Time-dependent induction of strand breaks upon exposure of plasmid DNA to 10 mM of ChOOH in the presence of Cu^2+

In order to better assess the DNA damage caused by ChOOH-derived oxidation products, time course experiments were carried out using 10 mM of ChOOH. As showed in Figures 3A₁ and B, the strand breaks increase with the time of DNA exposure. BER enzymes treatment follows the same time-dependent pattern, as showed upon incubation of the reactions with Fpg (Figure $3A_2$) or Endo III (Figure $3A_3$). However, the extension of damage is quite different. At all times of reaction, post-treatment with FPG increased $\sim 20\%$ the pre-existing damage caused by attack on the ribose phosphate backbone, while the post-treatment with Endo III was able to increase only ~ 10% the pre-existing damage. Taken together, these results show that ChOOH-decomposition products lead to base oxidations, preferentially at Fpgsensitive lesions.

Determination of 8-oxo-7,8-dihydro-2'-deoxyguanosine

Since 8-oxodGuo is the most common lesion excised by Fpg, it is worthy to confirm if ChOOH were able

to provoke this lesion. In this way, experiments using isolated dGuo and HPLC/MS/MS techniques were carried out. A typical chromatogram obtained after 2 h of ChOOH incubation with dGuo in the presence of 100 µM of copper ions is shown in Figure 4. Controls were made replacing the solution of ChOOH by ethanol. In the control incubation (Figure $4A_1$), a small amount of 8-oxodGuo was produced (area = 401). However, after 2 h of exposure to ChOOH and copper ions, the amount of 8-oxodGuo increased significantly (Figure $4B_1$, area = 3 819). To ensure correct measurements, an isotopically labelled 8-oxodGuo was added to the samples as an internal standard. The amount of internal standard did not vary significantly among different incubations (Figures 4A₂ and B_2). Similar reactions and the respective controls were carried out for each selected time (0.5, 1, 2, 4)and 6 h). For each time, an 8-oxodGuo/8-oxodGuo-N¹⁵ ratio was calculated, dividing the 8-oxodGuo area by the area of the internal standard. These results are shown in Figure 5. 8-oxodGuo is produced in a time dependent-manner, reaching the maximum level after 4 h of incubation. On the other hand, control reactions did not show significant levels of this lesion. A decrease in 8-oxodGuo amount was obtained after 6 h of incubation. This result can be explained by the fact that formed 8-oxodGuo can be further oxidized by ChOOH-decomposition products, since 8-oxodGuo is far more easily oxidized than the parent dGuo [29]. For example, it has been shown that 8-oxodGuo is a better substrate for ${}^{1}O_{2}$

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Figure 3. Time-dependent induction of strand breaks upon exposure of pBSK-II plasmid DNA to 10 mM of ChOOH and 10 μ M of Cu²⁺. Reactions mixtures were incubated for different times at 37°C in phosphate buffer (pH 7.4, with 30% ethanol, to allow ChOOH solubilization). Subsequently, ChOOH were extracted and DNA was subjected to a second incubation with DNA glycosylases for 30 min at 37°C. (A) Agarose gel electrophoresis. (1) Incubation in the absence of enzymes; (2) Incubation in the presence of 0.8 U of Fpg; (3) Incubation in the presence of 1 U of Endo III. Lanes 1–5: incubation for 0.5, 1, 2, 4 and 6 h, respectively. Plasmid DNA forms: OC: open circular; SC: supercoiled: L: linear. (B) Control percentage of supercoiled DNA. The SC percentage was calculated considering the SC amount of control incubation as 100%. Results represent the average and SD of at least three independent determinations.



Figure 4. Typical chromatograms for the detection of 8-oxodGuo by HPLC coupled to tandem mass spectrometry in the SRM mode. Reactions mixtures were incubated for 2 h at 37°C under agitation. After that, [M+5] isotopically labelled 8-oxodGuo was added to the samples and the ChOOH were extracted from the reactions prior to injection onto HPLC/MS/MS system. (A) Control chromatograms replacing ChOOH by ethanol. (B) Incubation of 10 mM of ChOOH with 5 mM of dGuo in the presence of 100 μ M of Cu²⁺. (1) Specific mass transition for 8-oxodGuo, 284 \rightarrow 168. (2) Specific mass transition for the internal standard, [M+5] isotopically labelled 8-oxodGuo, 289 \rightarrow 173.

oxidation, as compared with the parent dGuo [29,30]. In the same way, the 8-oxodGuo produced in ChOOH incubations can be consumed in further oxidative reactions and the measured amount can be a balance between the amount of produced and consumed 8-oxodGuo.

Discussion

Lipid peroxidation products have been implicated in DNA damage generation. For example, linoleic acid hydroperoxides were shown to cause single and double strand breaks in supercoiled plasmid DNA in the presence of iron ions [11]. In addition, arachidonic acid peroxidation was shown to induce DNA damage and mutations [12]. Additional evidences on the role of lipid hydroperoxides in causing DNA damage were obtained in this work with ChOOH. Copper catalysed decomposition of these hydroperoxides lead to plasmid DNA strand breaks in time and concentration-dependent manners.

It has been shown that copper ions occur in the mammalian cell nucleus [31,32]. This metal ion can be bound to the phosphate groups of the DNA backbone and to certain amino acids of some proteins, including chromatin [33–36]. Indeed, decomposition of organic peroxides by transition metals can generate peroxyl and alkoxyl radicals (showed in equations (1) and (2), respectively, as Ch-derived peroxyl (ChOO') and alkoxyl (ChO') radicals) [37]. An alternative mechanism, involving one-electron reduction of lipid hydroperoxides by Cu^{2+} , yielding alkoxyl radicals and Cu^{3+} , has also been proposed for alkoxyl radicals generation (equation 3) [38].



Figure 5. Time-dependent production of 8-oxodGuo upon exposure of 5 mM of dGuo to 10 mM of ChOOH and 100 μ M of Cu²⁺. Reaction mixtures were incubated for 0.5, 1, 2, 4 and 6 h at 37°C under agitation. After that, [M+5] isotopically labelled 8-oxodGuo was added to the samples and the ChOOH were extracted from the reactions prior to injection onto HPLC/MS/MS system. Control reactions were done replacing ChOOH for ethanol. For each time, the 8-oxodGuo/8-oxodGuo-N¹⁵ ratio was calculated dividing the 8-oxodGuo area by the area of the internal standard. Results represent the average and SD of at least three independent determinations.

$$\begin{array}{l} \text{ChOOH} + \text{Cu}^{2+} \rightarrow \text{ChOO}^{\bullet} + \text{Cu}^{1+} + \text{H}^+ (1) \\ \text{ChOOH} + \text{Cu}^{1+} \rightarrow \text{ChO}^{\bullet} + \text{HO}^- + \text{Cu}^{2+} (2) \\ \text{ChOOH} + \text{Cu}^{2+} \rightarrow \text{ChO}^{\bullet} + \text{HO}^- + \text{Cu}^{3+} (3) \end{array}$$

Furthermore, not only metal ions can be responsible for lipid-derived radicals formation. Many endogenous oxidative processes such as lipid peroxidation and protein oxidation may give rise to peroxyl radicals. Peroxyl radicals can attack a DNA molecule remote from their site of formation due to their relatively long lifetime when compared with other radicals [39]. The presence of lipid peroxyl radicals has been related to enhanced levels of 8-oxodGuo in nuclear and mitochondrial DNA [40].

Cholesterol-derived radicals can generate oxidative base modifications no matter their source. In addition, an initial damage on the nucleobases can also lead to secondary DNA strand scission [41]. The preferred pathway of DNA oxidation by ChOOH-decomposition products was base modifications, as evaluated by incubation of plasmid DNA with BER enzymes. As already mentioned, Fpg excises modified purines, while Endo III excises modified pyrimidines. Purines were shown to be favoured sites of ChOOH-derived products attack. For comparison, alkoxyl radicals, formed by photodecomposition of [4-(tert-Butyldioxycarbonyl) benzyl]triethylammonium chloride also generates a high number of Fpg-sensitive modifications and only a small amount of Endo III-sensitive lesions. Additionally, peroxyl radicals generated in arachidonic acid peroxidation were shown to preferentially oxidize guanine base and, to a minor extent, cytosine and thymine. The same study also presents

evidence for the involvement of 8-oxodGuo in the oxidation of DNA by arachidonic peroxyl radicals [13]. The results obtained in our work and also in previous ones are not surprising, considering the fact that dGuo is the most easily oxidizable nucleic acid base by ${}^{1}O_{2}$ and one-electron transfer mechanisms in DNA [42,43]. In this way, with the exception of the unselective hydroxyl radicals, other oxygen-derived radicals react preferentially with dGuo [44].

Korytowski et al. [45] have shown that different ChOOH isomers have different ability to induce free radical lipid peroxidation in cell membranes. These authors showed that, contrasting with the results obtained for 7 α -OOH and 5 α -OOH, the 6 β -OOH isomer was unable to initiate chain peroxidation, although all isomers were reduced by a lipophilic iron chelate and ascorbate (Fe(HQ)₃/AH⁻) at the same rate [46]. In the same way, the contribution of each ChOOH isomer in causing the overall oxidative DNA damage is now under investigation.

Intra-cellularly, LOOH might move for example, from mitochondria (which are under relatively high oxidative pressure) to the nucleus, where DNA could be damaged [47]. Phospholipids and unesterified Ch are known to move from one membrane compartment to another within cells. However, it has been shown that phospholipids translocate extremely slowly on their own (i.e. without protein mediation), whereas Ch does so relatively rapidly [48]. Indeed, phospholipid hydroperoxides translocate much more rapidly than their respective parent phospholipids [49]. The same behaviour is found for ChOOH, with rate constants for ChOOH transfer far exceeding that of parent Ch [14,47].

An interesting point that should be mentioned is the unexpected high levels of oxysterols observed in human plasma, LDL and erythrocytes. In contrast to polyunsaturated fatty acids, Ch does not contain bisallylic hydrogen and the reactivity of allylic hydrogen toward peroxyl radical, a chain-carrying species in lipid peroxidation, is ~ 100 times smaller than that of bisallylic hydrogen. However, the high level of oxygenated oxysterols may be ascribed to slower metabolism and/or excretion [50]. At least three different classes of intracellular enzymes have been implicated in reductive LOOH detoxification: GSH peroxidases [51], peroxiredoxins [52] and GSH-S-transferase [53]. The former class includes the enzymes GPx-1 and GPx-4 and is considered to be more important in overall LOOH disposal [2]. However, GPx-1 cannot directly act on phospholipid hydroperoxides in membranes, unless sn-2 fatty acyl bonds are first hydrolysed to liberate the peroxidized fatty acids. Only GPx-4 has this ability [51,54]. More interestingly, ChOOH are completely resistant to GPx-1 detoxification and are also more resistant to GPx-4-dependent elimination than fatty acid hydroperoxides and their esterified form [15,55].

Cholesterol is found in the membrane of all eukaryotic cells. The fact that nuclear membranes contain much less Ch than plasma membranes $(\sim 10\%$ in nuclear membranes when comparing with 40% of the total lipid of plasma membranes) can be overwhelmed by the relatively rapidly translocation of ChOOH among membranes, along with their slower detoxification rates [2,56]. Moreover, oxidation of nuclear membrane cholesterol was shown to inhibit the activity of nucleoside triphosphatase (NTPase), an enzyme believed to be involved in nucleo-cytoplasmic trafficking [57]. Unspecified lipid peroxidation products were also implicated in nuclear effects. For example, it was shown that primary photosensitization reactions involving ¹O₂ generation in membranes lead to DNA damage. This effect was probably caused by secondary reactive species derived from lipid peroxidation [58]. Therefore, ChOOH should be considered as potential effectors that can play a role in DNA oxidative damage.

In summary, this work showed that ChOOHderived products oxidize DNA, especially at Fpglabile sites. The results obtained with isolated dGuo and HPLC/MS/MS analyses support the dGuo oxidation in plasmid DNA. Alkoxyl and/or peroxyl radicals derived from reaction of ChOOH with copper ions should be responsible for the damage. DNA damage caused by putative hydroxyl radicals derived from secondary reactions can be ruled out, since experiments were performed employing 30% ethanol as co-solvent. Ethanol is a well known scavenger of hydroxyl radicals [3].

Despite the recognized role of catalytic metals in inducing lipid peroxidation, the exact mechanism is still not clear. It has been argued from thermodynamic calculations that LOOH are unable to reduce Cu^{2+} to Cu^{1+} , and only the last one interacts with LOOH [59]. However, another study showed that in the presence of an oxidizable substrate, LOOH are capable of reducing Cu^{2+} yielding Cu^{1+} and lipid-derived peroxyl radicals [60]. The last work proposed a reversible intermediate complex, denoted as [Cu^{2+} -LOOH], which will generate Cu^{1+} and peroxyl radical.

In this view, DNA damage generated by ChOOH in the presence of copper ions can be started by the reaction of ChOOH and Cu^{2+} , yielding ChOO[•] and Cu^{1+} (Figure 6). Besides the equilibrium of equation (1) lies to the left, in the presence of an oxidizable substrate (in this case, DNA molecule) the thermodynamic barrier can be overcome, and ChOO[•] and Cu^{1+} will be formed. Then, a redox cycling can be proposed. Cu^{1+} ions react with another ChOOH molecule, yielding ChO[•] and regenerating Cu^{2+} . The ChO[•] are more reactive than ChOO[•], being able to react with DNA. Also, ChO[•] can undergo cyclization,



Figure 6. Possible pathways for ChOOH-derived DNA damage.

vielding carbon-centred epoxyallylic radicals (OCh[•]). The later radicals can couple with molecular oxygen, forming epoxyperoxyl radicals (OChOO'), further amplifying the damaging agents able to react with DNA [61]. Another possibility for ChO[•] is β -scission reaction forming aldehydes and alkyl radicals (Ch[•]), which also couple with O₂, yielding ChOO'. Additionally, self-termination of peroxyl radicals will give rise to ¹O₂, as proposed by Russell [62] and confirmed for lipid peroxides in the presence of metal ions or other oxidants [63-65]. Singlet molecular oxygen can react with dGuo base, yielding 8-oxodGuo [42]. The initial reaction of ChOOH with Cu^{2+} , vielding ChO[•] and Cu³⁺, must also be considered [38]. Expected cholesterol end-products are 5,6epoxide, 7-ketone and the diols derived from decomposition of each hydroperoxide, especially the stable 7-OH [46,54,66].

ChOOH-derived DNA damage adds further support to the role of lipid peroxidation in inducing DNA modifications and mutation. Moreover, ChOOH can be considered to play a special role in the development of pathological conditions linked to DNA damage, giving their role in translocation through membranes, disseminating the peroxidative stress [67]. The results obtained here point to the importance of ChOOH as damaging agents to DNA and warranty further studies about the involvement of these species in the aetiology of some diseases.

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Declaration of interest

The authors report no conflicts of interest. The authors alone are responsible for the content and writing of the paper.

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