DNA repair phenotype and dietary antioxidant supplementation

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Phytochemicals may protect cellular DNA by direct antioxidant effect or modulation of the DNA repair activity. We investigated the repair activity towards oxidised DNA in human mononuclear blood cells (MNBC) in two placebo-controlled antioxidant intervention studies as follows: (1) well-nourished subjects who ingested 600 g fruits and vegetables, or tablets containing the equivalent amount of vitamins and minerals, for 24 d: (2) poorly nourished male smokers who ingested 500 mg vitamin C/d as slow- or plain-release formulations together with 182 mg vitamin E/d for 4 weeks. The mean baseline levels of DNA repair incisions were 65.2 (95% CI 60.4, 70.0) and 86.1 (95% CI 76.2, 95.9) among the male smokers and well-nourished subjects, respectively. The male smokers also had high baseline levels of oxidised guanines in MNBC. After supplementation, only the male smokers supplemented with slow-release vitamin C tablets had increased DNA repair activity (27 (95% CI 12, 41) % higher incision activity). These subjects also benefited from the supplementation by reduced levels of oxidised guanines in MNBC. In conclusion, nutritional status, DNA repair activity and DNA damage are linked, and beneficial effects of antioxidants might only be observed among poorly nourished subjects with high levels of oxidised DNA damage and low repair activity.

DNA repair: Oxidatively damaged DNA: Phytochemicals: Vitamin C: Vitamin E

Cellular DNA may be protected against oxidation by antioxidants, and oxidised DNA lesions are removed by several repair systems such as base excision repair, nucleotide excision repair and mismatch repair that have overlapping specificity and may interact or function as back-up systems. Low activity of DNA repair, towards oxidised DNA may increase the risk of cancer by increasing the accumulation of errors in the genome. It has been reported that low repair activity towards oxidised DNA is associated with increased risk of cancer in case–control studies, although this could be due to reverse causality. The assessment of DNA damage has been widely discussed and evaluated in dietary intervention studies, whereas assays for the assessment of DNA repair are few and data are limited.[3]

In principle, there are two different approaches to measure DNA repair activity in cells. Lesions can be introduced in cellular DNA and the repair activity is assessed as the removal of lesions during a subsequent incubation period. The most popular type of repair activity measured by the single cell gel electrophoresis (comet) assay has been the measurement of rejoining of strand breaks after ex vivo exposure to clastogens such as H₂O₂ or ionising radiation. Although these assays for DNA repair activity are simple, they suffer from the drawback that the genotoxic exposure may damage the repair proteins and initiate the process of programmed cell death (apoptosis). Moreover, strand breaks are generated in cultured lymphocytes as a result of exposure to atmospheric O₂ and they can also accumulate as a consequence of poor repair activity. The validity of repair assays based on the rejoining of strand breaks in antioxidant intervention studies has been questioned because they cannot distinguish between rejoining of strand breaks and ex vivo scavenging effects by antioxidants.[7] These problems can be overcome by assays where extracts of cells are incubated with substrate DNA containing a defined number of lesions. Recently, a novel version of the comet assay was developed for the measurement of oxoguanine repair activity of cell extracts using substrate nuclei treated with the Ro19-8022 photosensitiser and white light.[6] This photosensitiser generates 8-oxo-7,8-dihydro-2′-deoxyguanosine (8-oxodG) and very few strand breaks.[8] The repair activity is measured as the number of incisions that are generated during the incubation. This approach has been used in a few studies and indicates that dietary antioxidants can alter DNA repair activity.[9–11] The glycosylase enzymes incise at the sites of oxidised bases, leaving apurinic/apyrimidinic sites that are converted into breaks in the assay. The repair activity of Ro19-8022-damaged DNA lesions is assessed as the removal of DNA incisions.

Abbreviations: FPG, formamidopyrimidine DNA glycosylase; MNBC, mononuclear blood cells; OGG1, oxoguanine glycosylase 1; 8-oxodG, 8-oxo-7,8-dihydro-2′-deoxyguanosine.

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DNA is believed to mainly represent oxoguanine glycosylase 1 (OGG1), whereas nei-endonuclease VIII-like (for example, NEIL1) glycosylase and nucleotide excision repair enzymes probably have little contribution to the overall repair phenotype. The aim of the present study was to investigate if fruit and vegetable intake or antioxidant supplementation modulate the oxoguanine repair activity in mononuclear blood cells (MNBC) of human subjects. The reliability of the repair assay was assessed in liver homogenate tissues of wild-type and Ogg1 knockout mice.

Materials and methods

We investigated the DNA repair activity of cryopreserved MNBC from two antioxidant intervention studies that have been described in detail previously12–14. The characteristics of the subjects enrolled in the two studies are outlined in Table 1. The level of oxidised purines was measured as formamidopyrimidine DNA glycosylase (FPG)-sensitive sites by the comet assay; these lesions are mainly represented by 8-oxodG and ring-opened purine lesions.

Characteristics of study 1 – ‘six-a-day’ study

The ‘six-a-day’ study was a parallel placebo-controlled intervention study in which forty-three subjects were randomised into three groups ingesting an energy-balanced basal diet free from fruits, vegetables and antioxidants, and supplemented with (1) 600 g fruits and vegetables, (2) tablets containing the corresponding amounts of vitamins and minerals, or (3) placebo tablets. The study is best characterised as a combined depletion and supplementation study. Blood and urine samples were collected at baseline (before) and at days 9, 16 and 24 of the supplementation. Post-intervention samples were obtained 4 weeks after the intervention had ceased. The constituents of the fruit and vegetable diet and content of the tablets have been described in detail previously12,13,14. The samples analysed in the present study were obtained at baseline (pre-intervention), end-supplementation (day 24) and post-supplementation (day 53).

<table>
<thead>
<tr>
<th>Study...</th>
<th>’Six-a-day’</th>
<th>Vitamin C and E</th>
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</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>sd</td>
</tr>
<tr>
<td>Age (years)</td>
<td>27</td>
<td>6</td>
</tr>
<tr>
<td>Sex (n)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Men</td>
<td>22</td>
<td></td>
</tr>
<tr>
<td>Women</td>
<td>21</td>
<td></td>
</tr>
<tr>
<td>Alcohol intake (g/d)</td>
<td>15</td>
<td>18</td>
</tr>
<tr>
<td>Smoking</td>
<td>Non-smokers</td>
<td></td>
</tr>
<tr>
<td>Baseline vitamin C (µmol/l)</td>
<td>69</td>
<td>17</td>
</tr>
<tr>
<td>Baseline level of FPG sites (arbitrary units)</td>
<td>37†</td>
<td>24</td>
</tr>
</tbody>
</table>

FPG, formamidopyrimidine DNA glycosylase.

The subjects enrolled in the study had a high baseline plasma concentration of vitamin C. The concentration of plasma vitamin C decreased dramatically in the placebo group (the mean concentration of vitamin C was 21 µmol/l at the end of the supplementation). The profile of carotenoids was somewhat more complex; the plasma lycopene concentration only increased in the group that ingested 600 g fruits and vegetables/d, whereas the plasma β-carotene concentration increased the most in the group of subjects receiving tablets14.

Characteristics of study 2 – vitamin C and E study

The vitamin C and E study was designed as parallel, blinded, placebo-controlled investigation with forty-eight male smokers randomised into three groups as follows: (1) tablets with plain-release formulation of vitamin C (2 x 250 mg/d) and vitamin E (2 x 91 mg/d); (2) tablets with slow-release formulation of vitamin C (2 x 250 mg/d) and vitamin E (2 x 91 mg/d); (3) placebo. Whereas the content of vitamin C in the tablets is identical, ingestion of the tablets yields pharmacokinetically different variables due to the differences in release of vitamin C (for example, there is less fluctuation of the plasma concentration by ingestion of the tablets with slow-release formulation of vitamin C as compared with the plain-release tablets). On the contrary, the different formulations of the tablets have little pharmacokinetic effect on the plasma concentration of vitamin E15. The subjects ingested tablets daily for 4 weeks. The samples analysed in the present study were obtained at baseline (pre-supplementation) and at the end of the supplementation (end-supplementation).

Measurement of DNA repair activity

The validation of the repair assay was assessed in liver extracts from wild-type and Ogg1−/− mice. Afterwards, the DNA repair activity was determined in MNBC of the two intervention studies. Assessment of DNA repair incisions was analysed by the comet assay as described by Collins et al.6, with modifications as described below.

Preparation of cells with substrate nuclei

We used A549 lung epithelial cells as substrate nuclei (American Type Culture Collection, Manassas, VI, USA). These were washed twice with PBS before treatment. The A549 cells were treated with a 1 µM solution of the Ro19-8022 photosensitiser (a gift from Hoffmann-La Roche, Basel, Switzerland) dissolved in PBS, and irradiated for 4 min at 33 cm from two 500 W halogen lamps. The irradiation of the cells suspension was carried out on ice to avoid heating of the solution. After the treatment, cells were centrifuged at 700 g for 15 min to remove the photosensitiser, washed in RPMI-1640 medium (Gibco RBL, Grand Island, NY, USA) and centrifuged at 400 g for 15 min. The pellet was re-suspended in freezing medium (50% fetal bovine serum (Gibco RBL), 40% RPMI-1640 medium and 10% dimethyl sulfoxide) to a concentration of 3 x 10^6 cells/ml and cryopreserved in small samples at −80°C.
Preparation of mouse liver extract

Extracts of livers were obtained by placing the tissue into a stainless-steel cylinder and by use of a plunger forcing it through a sieve in one end of the cylinder (0.5 cm diameter, mesh size 0.4 mm), while the cylinder was submerged in 2 ml ice-cold buffer A (45 mM-HEPES, 0.4 M-KCl, 1 mM-EDTA, 0.1 mM-dithiothreitol, 10 % glycerol, adjusted with KOH to pH 7.8) according to the procedure described by Møller et al.16 Subsequently, the extracts were cryopreserved at −80°C. The Ogg1−/− mouse liver was a cryopreserved sample used in a previous study17. On the day of the experiment, samples of the cryopreserved samples were thawed and processed as described for MNBC.

Preparation of cell extract from human mononuclear blood cells

The same isolation procedure of MNBC was used in the ‘six-a-day’ and vitamin C and E studies. MNBC from venous blood were isolated in control preparation tubes and cryopreserved in freezing medium (50 % fetal bovine serum, 40 % RPMI-1640 medium and 10 % dimethylsulfoxide). The cryopreserved cell samples were thawed and 5 ml of a 3-fold diluted extraction buffer A were added. The number of cells was ascertained and the samples centrifuged at 700 g for 5 min at 4°C. As much as possible of the supernatant fraction was removed. The pellet was re-suspended by vigorously tapping the tube and 20 μl buffer A was added for each 106 cells. The suspended cells were divided into 50 μl samples and frozen at −80°C. This procedure of MNBC extract generation deviated from the original procedure6, but we obtained similar incision activity in control experiments where we compared the two extract isolation procedures (results not shown).

In vitro repair incubation

Frozen samples of MNBC extracts were thawed and 12 μl 1 % Triton X-100 in buffer A were added. The lysate was subsequently centrifuged at 13 500 rpm for 5 min at 4°C to remove cell debris. The supernatant fraction was mixed with 4 volumes buffer B (40 mM-HEPES, 0.1 M-KCl, 0.5 mM-EDTA, bovine serum albumin (0.2 mg/ml), pH 8) and kept on ice until use. Samples of cells with substrate nuclei were thawed, mixed with low-melting-point gel (0.75 %; Sigma-Aldrich, Brondby, Denmark) and applied on 85 × 100 mm GelBond® films (Cambrex, Medinova Scientific A/S, Hellerup, Denmark). Each GelBond consisted of eight 19 × 23 mm agarose gels. The GelBonds were immersed in lysis solution (2.5 mM-NaCl, 0.1 mM-Na2EDTA, 10 mM-tri(hydroxymethyl)-aminomethane (Tris), 1 % Triton X-100, pH 10) for 1 h at 4°C. Then the GelBonds were washed three times (5 min each) in buffer B. Cell extract (60 μl) was added to each gel and incubated for 20 min at 37°C in a humid box. Control gels were incubated for 20 min with 60 μl of a control solution consisting of Triton X-100 and buffer B (without lymphocyte extract). The GelBonds were then placed in a horizontal electrophoresis tank and immersed in fresh solution (0.3 M-NaOH, 1 mM-Na2EDTA) for 40 min, before electrophoresis at 25 V (0.83 V/cm) and 300 mA for 20 min. After neutralisation with 0.4 M-Tris-HCl (pH 7.5) cells were placed in 96 % ethanol for 1.5 h or overnight. The nuclei were visualised in an Olympus fluorescence microscope at 40 × magnification and two gels of 100 nuclei were scored as degrees of migration in classes 0–4 after staining with 50 μl YOYO-1 (Molecular Probes, Leiden, The Netherlands) in PBS buffer. The repair activity of the MNBC extract was determined as the difference in score (arbitrary units) between parallel gels incubated with extract and control solution.

Statistics

Differences in the baseline values of DNA repair phenotype of the two datasets were tested by ANOVA with unequal variance of groups because of lack of homogeneity of the variance (Levene’s test). Effects of the intervention were tested by one-factor repeated-measurements ANOVA of two groups (placebo group v. active group). Since the data from the placebo groups were tested twice against the active groups, the statistical significance was accepted at P<0.025 after Bonferroni correction for multiple tests. Analysis of normal distribution of residuals was tested by the Shapiro–Wilk W test. The relationship between DNA repair incision activity and FPG sites in baseline samples was analysed by linear regression with P<0.05 as the significance level. The statistical analysis was carried out in STATISTICA version 5.5 for Windows (Statsoft, Inc., Tulsa, OK, USA).

Results

The DNA repair assay was implemented with several modifications compared with the original protocol, but we essentially observed similar time curves of the incision activity as reported previously, i.e. an initial linear increase of incisions followed by a plateau6. We found that the most reliable incision activity was observed after 20 min incubations, based on considerations that the number of incisions should be as high as possible before the plateau is reached (results not shown). Fig. 1 outlines the results from experiments of dilutions of enzyme extract from the liver of mice. The protein concentration of the undiluted liver extract was 4 mg/ml, which was lower than that used for MNBC (about 20 mg/ml) because we assumed higher repair activity of the hepatocytes compared with MNBC. The 10-fold dilution of the wild-type liver extract reduced the incision activity to 43 (95 % CI
21, 65\%). The repair incision activity of the \(Ogg1^{-/-}\) mouse liver extract was very low, i.e. 17 (95 \% CI 0, 39) \% compared with the undiluted wild-type mouse liver sample, but it was not significantly lower than the 100-fold and 1000-fold diluted wild-type mouse liver extract samples.

The results of the baseline values in the two intervention studies are shown in Fig. 2. As can be seen, the distributions are quite different \((P < 0.001; \text{Levene’s test})\). The values of the poorly nourished smokers in the vitamin C and E study are less scattered than the results from well-nourished subjects in the ‘six-a-day’ study, although there was difference in repair incisions between the placebo and slow-release groups at baseline \(( P < 0.05; \text{one-factor ANOVA})\). The mean repair incision activity was higher for the subjects in the ‘six-a-day’ study (mean 86·1; 95 \% CI 76·2, 99·9) than in the vitamin C and E study (mean 65·2; 95 \% CI 60·4, 70·0; \( P < 0.001; \text{ANOVA for groups with unequal variances})\). The residuals of this analysis did not deviate from a normal distribution \((P = 0.12; \text{Shapiro–Wilk W test})\), indicating little unexplained variation. The subjects in the vitamin C and E study were smokers, who had higher alcohol consumption, lower plasma vitamin C concentration, and higher levels of FPG sites in MNBC at baseline (Table 1). Although, at group level, there appeared to be an inverse relationship between the level of DNA repair incisions and FPG sites, there was not a statistically significant linear relationship between these biomarkers in the ‘six-a-day’ and vitamin C and E studies \(( P > 0.05; \text{linear regression analysis})\).

Table 2 outlines the results of repair phenotype of MNBC extracts. The analysis of DNA repair phenotype was successful in forty-six samples of the vitamin C and E study, encompassing eighteen datasets in each of the active supplementation groups and ten subjects in the placebo group. In the ‘six-a-day’ study, samples from thirty-nine subjects were available and analysed, which included twelve, thirteen and fourteen subjects in the vitamins, fruit and vegetable, and placebo group, respectively. There was no significant variation in the DNA repair phenotype related to the intervention \(( P > 0.025; \text{ANOVA for repeated measurements})\).

On the contrary, ingestion of slow-release vitamin C formulation for 4 weeks in heavy smokers was associated with a significantly increased repair capacity \(( P < 0.025; \text{repeated-measurement ANOVA})\), whereas there was no difference in DNA repair incisions between subjects supplemented with tablets with plain-release formulation of vitamin C and placebo \(( P > 0.025; \text{repeated-measurement ANOVA})\). In Fig. 3, changes in DNA repair activity observed in the present study are combined with changes in FPG sites previously reported\(^1^3\). The alteration of FPG sites is virtually a mirror image of the change in DNA repair.

**Discussion**

Oxidised DNA in cells is usually kept at a low steady-state level because the DNA repair system removes lesions from the DNA. However, during oxidative stress, the level of oxidised DNA in cells may increase because the DNA repair system cannot remove lesions with sufficient speed. Lower levels of oxidised DNA following supplementation with phytochemicals suggest that the bioactive constituents may increase the activity of the DNA repair system in addition to a direct scavenging effect of reactive oxygen species.

Methods for the assessment of DNA repair activity are scarce despite their relevance for understanding the cellular response to oxidative stress. The novel application of the comet assay for repair incision activity of cellular extracts is still relatively new and only a few groups have reported results from investigations with this method. Therefore we devoted some time to reassessing the initial assay validation carried out by Collins and co-workers, who reported that fibroblast cell cultures from \(Ogg1^{-/-}\) mice had very little incision activity\(^6\). We found a dramatically reduced incision activity of the diluted wild-type extract, and 10-fold dilutions did not differ significantly from \(Ogg1^{-/-}\) liver extract, which had virtually no incision activity on Ro19-8022 substrate nucleoids. These results are in accordance with those reported by Collins et al.\(^9\) who showed virtually no incisions by four-fold dilution of lymphocyte extracts.

Reliable assays for the measurement of repair of oxidised DNA bases by human MNBC cell extracts have become increasingly popular in recent years. However, investigations have mainly been carried out in small groups of subjects and high inter-individual variation can be difficult to separate from assay variation. In the present study we investigated the repair activity of MNBC extracts in two populations with different characteristics. This showed that at baseline there was not only a difference between subjects, but also differences between the groups of the population in the level of repair activity. Baseline differences in effect markers can be observed in parallel designs because of the unintentional heterogeneous randomisation of subjects. We do not believe that it affects the interpretation of the data because we have not selected participants with low incision activity that otherwise could lead to a regression towards the mean phenomenon. In addition, the data obtained in the vitamin C and E study are not unusually low; for example, we have previously found incision activities in the range of 50 arbitrary units in young and healthy subjects\(^1^8\). However, problems with baseline differences in biomarkers can be alleviated by the use of cross-over designs, but this type of design suffers from other problems such as carry-over effects.

![Fig. 2. Distributions of baseline DNA repair incision of mononuclear blood cell of subjects in the ‘six-a-day’ (■, –) and vitamin C and E (○, -) intervention studies. The data are repair incisions measured in arbitrary units.](image-url)
and systematic variation over time can be particularly troublesome. The overall mean of baseline repair activity for both datasets was 76.7 (SD 29.6) with a corresponding CV of 39%, but the data are not normally distributed. This is similar to the CV (34%) reported in a study on 375 subjects\(^{19}\), whereas another study among 244 subjects reported a CV in the range of 100%\(^{20}\). The repair incision activity in these subjects was reported to correlate weakly and positively with age\(^{19}\) or not being age dependent\(^{20}\). Sex and smoking habit were not reported to affect DNA repair activity towards oxidised bases, whereas polymorphisms in the \(OGG1\) and \(XPA\) genes affected the repair incision activity\(^{19,20}\). This indicates that age, sex and smoking are not strong determinants for the repair incision activity and the differences we have observed between the subjects in the ‘six-a-day’ and vitamin C and E studies could be due to other lifestyle and genetic variables. It should be emphasised that the repair activity of the two datasets were analysed at the same time, and the difference cannot be attributed to variation in the analysis over time. The ‘six-a-day’ samples had been stored for a longer time (6 years) than the samples from the vitamin C and E study (4 years). A recent study did not show decreased repair activity, determined as cleavage of 8-oxoG-containing oligonucleotides, in samples after storage for 3 years\(^{21}\). Clearly, further studies are warranted that assess the stability of repair activity during storage.

The effect of phytochemical supplementation on DNA damage has been investigated in more than 100 studies, whereas few studies on repair activity are available\(^4\). Increased DNA repair activity was observed after supplementation with kiwi fruits, and this was accompanied by decreased levels of oxidised DNA in MNBC\(^{10}\). Using a different assay for repair activity, cell extract-associated incision activity of plasmid DNA treated with

### Table 2. DNA repair incisions in mononuclear blood cell extracts of subjects in the ‘six-a-day’ and vitamin C and E studies* (Mean values with their standard errors)

<table>
<thead>
<tr>
<th>Study</th>
<th>Pre-supplement</th>
<th>End-supplement</th>
<th>Post-supplement</th>
<th>P†</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Mean</td>
<td>SEM</td>
<td>Mean</td>
<td>SEM</td>
</tr>
<tr>
<td><strong>Six-a-day</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Fruit and vegetables (n 13)</td>
<td>88.1</td>
<td>9.6</td>
<td>82.4</td>
<td>8.9</td>
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<tr>
<td>Vitamins (n 12)</td>
<td>77.1</td>
<td>14.2</td>
<td>78.6</td>
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<tr>
<td>Placebo (n 14)</td>
<td>97.4</td>
<td>11.5</td>
<td>91.7</td>
<td>11.1</td>
</tr>
<tr>
<td><strong>Vitamin C and E</strong></td>
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<td></td>
</tr>
<tr>
<td>Vitamin C (slow-release formulation) and vitamin E (n 18)</td>
<td>58.3</td>
<td>3.3</td>
<td>74.0</td>
<td>5.3</td>
</tr>
<tr>
<td>Vitamin C (plain-release formulation) and vitamin E (n 18)</td>
<td>67.7</td>
<td>3.8</td>
<td>72.6</td>
<td>4.6</td>
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<td>Placebo (n 10)</td>
<td>73.0</td>
<td>5.1</td>
<td>71.6</td>
<td>4.6</td>
</tr>
</tbody>
</table>

* DNA repair incisions are obtained as the difference in score (arbitrary units) of substrate samples (i.e. A549 cells exposed to 1 \(\mu\)M-Ro19-8022 and white light) treated with mononuclear blood cell extract and buffer. The \(P\) values correspond to statistical analysis of repeated-measurement analysis that tests the difference in changes over time between the treatment and placebo groups.

† Effects of the intervention were tested by one-factor repeated-measurements ANOVA with \(P\leq0.025\) as the level of statistically significant effect.
methylene blue and visible light, it was indicated that ingestion of cooked carrots increased repair activity, whereas ingestion of mixed carotene tablets and tinned mandarin oranges had no effect on DNA repair activity. Increased repair incisions of Ro19-8022-induced DNA damage were also observed following a 1-week supplementation with coenzyme Q10, whereas the levels of FPG sites in MNBC did not change. Our observation that increased incision repair activity was induced by vitamin C supplementation corroborates well with these results, and indicates that it is possible to increase repair activity towards oxidised DNA in MNBC by supplementation with phytochemicals and antioxidants. It is all the more interesting that it was not possible to alter the repair incision activity in MNBC of the well-nourished subjects in the ‘six-a-day’ study. Measurements of the OGG1 mRNA levels in whole blood of the subjects were not altered because of the treatment in the present study, and the vitamin C and E supplementation study, although period effects were observed in both studies. However, the data on OGG1 mRNA levels are not directly comparable with the measurements of DNA repair activity because the latter was investigated in MNBC. The unaltered incision repair activity of the subjects in the ‘six-a-day’ study might be explained by the short duration of the depletion period in these well-nourished subjects. The lifespan of lymphocytes, which is the major fraction of MNBC, is usually considered to be long (probably years), although it is likely to depend on the subtype of lymphocytes. The depletion of phytochemicals in the ‘six-a-day’ study was only 24 d, which is a short time compared with the lifespan of lymphocytes. Thus, it is reasonable to expect that a large fraction of the cells in the blood remained unaltered after 24 d. A similar lack of effect was observed for antioxidant enzyme activity (including catalase, superoxide dismutase and glutathione reductase) in erythrocytes, which only have a lifespan of 120 d, but glutathione peroxidase in the erythrocytes was significantly increased by the fruit and vegetable intervention, indicating that changes in some biomarkers over a shorter time period may take place even in cell populations which have long half-lives even without de novo protein synthesis. It had been shown that the ex vivo sensitivity to oxidatively induced DNA damage increased in subjects with reduced antioxidant levels at baseline associated with a controlled run-in period with a diet poor in antioxidants. Taken together, the data suggest that the subjects in the ‘six-a-day’ study might not have experienced significant oxidative stress following the 24 d depletion of antioxidants. Thus, the most important interpretation of these observations is that swift alterations in DNA repair enzyme activity can be observed by supplementation of antioxidants in poorly nourished subjects, whereas severe depletion of antioxidants for long periods of time might be required to observe alterations in well-nourished subjects.

In conclusion, determinants of the oxoguanine glycosylase repair phenotype encompass dietary habits, lifestyle factors and genetic differences. Subjects with poor habitual antioxidant intake may benefit from supplementation with concomitant reduction of the level of oxidised DNA in cells.

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