Role of Oxidative DNA Damage and Antioxidative Enzymatic Defence Systems in Human Aging

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Abstract: Oxidative stress is regarded as a main causal factor for natural aging. This study tested the hypothesis that healthy elderly people show higher oxidative DNA damage levels and lower antioxidative enzymatic defense capacities than younger ones. In a cross-sectional study, blood samples of 20 older (62-79 years) and 20 younger adults (24-28 years) were compared with respect to oxidative DNA damage in lymphocytes (alkaline elution), oxidative status (serum peroxides), activities of superoxide dismutase (SOD) and glutathione peroxidase (GPx) and concentrations of total glutathione. In accordance with our hypothesis, elderly males showed a tendency towards higher levels of oxidative DNA damage (single strand breaks). SOD activity inversely correlated with the amount of DNA damage (single-strand breaks and Fpg-sensitive modifications). Oxidative status was increased in older men and negatively correlated with glutathione concentrations. GPx activity was elevated and the SOD/GPx ratio lowered in older males. Subjects with lowered SOD/GPx ratio showed increased oxidative DNA damage. The results indicate age-related changes in the balance between first step (SOD) and second step (GPx) of the enzymatic antioxidant defense system. They support the assumption that a biological optimum between antioxidative enzymes might be more important than their absolute activities.

Key Words: Aging, oxidative DNA damage, oxidative status, antioxidative enzymes, superoxide dismutase, glutathione peroxidase, glutathione.

INTRODUCTION

Different hypotheses have been proposed to explain the aging processes in humans. Among the related theories, the most advanced is the oxidative damage theory, which has been developed from the free radical theory of Harman [1, 2]. It considers oxidative stress as main causal factor for natural aging and is based on the high reactivity of reactive oxygen species (ROS), which are ubiquitously generated byproducts of the oxygen metabolism in living beings [3-5]. An imbalance between oxidative and reductive processes in favor of oxidative ones is called oxidative stress and supposed to result in extensive oxidative damage to nucleic acids, proteins and lipids [6-8]. These damaged molecules perturb cellular homeostasis, reducing more and more functionability and life span of cells and finally of organs [9]. Studies in animals indeed have indicated that oxidative damage accumulates during life in proteins, lipids and DNA [10]. In humans, an excess of ROS has been made responsible for premature aging and the development of atherosclerosis and cancer [11-13]. Increased oxidative stress has also been found in patients with chronic renal failure [14].

Oxidative stress is caused in humans by various pathophysiological mechanisms, in particular by the increased production of ROS during inflammatory processes and by various environmental factors (ionizing radiation, nicotine, because of its mispairing with adenine [21]. Increases of 8oxo-dG with age have been observed by several groups, e.g. in rats and human brain [22, 23]. A down-regulation of antioxidative protective mechanisms as well as an increased production of ROS might be important factors for an age-related increase of oxidative damages. Studies on murine aging have shown that the activity of SOD1 rose during aging in all organs studied, whereas the activities of GPx1 and CAT displayed organ-specific profiles [24]. Despite evidence for altered oxidative stress

alcohol, unbalanced diet, etc.). Protective enzymatic and non-enzymatic antioxidant defense mechanisms reduce oxi-

dative stress by degradating ROS. Major intracellular anti-

oxidative enzymes are the superoxide dismutase (SOD),

catalase (CAT), and glutathione peroxidase (GPx). They act

in two steps: firstly, SOD converts the highly active superox-

ide radicals (O_2) into hydrogen peroxide (H_2O_2) and oxygen

(O₂). Afterwards, CAT and GPx independently convert H₂O₂

to water and oxygen [15]. Glutathione (GSH) is not only a

cofactor for GPx, but can also react as a direct scavenger of

ROS [16]. On the non-enzymatic level, also vitamins (vita-

min C, vitamin E and β -carotene) and other antioxidant

compounds scavenge free radicals and delay oxidation of

molecules. While antioxidant defense mechanisms inhibit

damage generation, DNA repair enzymes (glycosylases)

remove DNA base modifications generated by ROS. One of

the most abundant DNA lesions resulting from reaction of

ROS with DNA is 7,8-dihydro-8-oxo-2'-deoxyguanosine (8-

oxo-dG), a modification caused by hydroxylation of the C-8

position of guanine [17-20]. It causes a $G \rightarrow T$ transversion

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levels and antioxidative enzyme activities with aging [25-31], there are only few data on the relationship between DNA damage and these parameters in healthy elderly people. Therefore, the objective of this study was to test the hypothesis that biological processes of aging are associated in humans with an increase of oxidative stress and a decrease of both antioxidative capacity and DNA repair mechanisms. For the first time, the complex method of alkaline elution was applied to healthy elderly for determining oxidative DNA damage.

SUBJECTS AND METHODOLOGY

Subjects

In a cross-sectional study design, 20 healthy younger adults (10 males, 10 females, age 24-28 years, mean age 25.8 years) were compared to 20 free-living healthy older adults (10 males, 10 females, 62-79 years, mean age 69.2 years) with respect to their oxidative status (serum peroxide level), oxidative DNA damage, antioxidative enzyme activities (SOD and GPx), total glutathione, and antioxidative micronutrients (B-carotene, vitamin C, and zinc). Since disease may influence plasma antioxidant measurements, exclusion criteria were the presence of inflammatory processes as well as acute and chronic diseases. Because nutrition and life style-related parameters may also influence antioxidative parameters, they were additionally assessed by questionnaire. The study protocol was approved by the Ethics Committee of the Medical Association of Rhineland-Palatinate, Germany. All volunteers signed written informed consent forms before entering the study.

Biochemical Methods

After an overnight fast, 30 ml peripheral venous blood was taken in the morning from each subject by a physician at the Department of Sports Medicine, University of Mainz. All blood samples of a subject were taken on the same day and analyzed within two days.

Oxidative DNA Damage

The level of oxidative DNA damage in blood lymphocytes was measured by the alkaline elution technique. The assay originally described by Kohn et al. [32] makes use of the fact that the elution rate of mammalian DNA from a membrane filter depends on the average length of the DNA molecules and therefore, the number of DNA single-strand breaks. Several other DNA modifications can be quantified with the same sensitivity, since they can be converted into single-strand breaks by a preincubation of the DNA with suitable repair enzymes. In this study, formamidopyrimidine N-glycosylase (Fpg), the bacterial functional homolog of the mammalian repair glycosylase OGG1, was used to quantify its substrate modifications (Fpg-sensitive modifications), which include 8-oxo-dG, formamidopyrimidines and sites of base loss (AP sites) [33]. The assay was carried out as described previously [34, 35]. In brief, 8 ml human venous blood was collected in citrate-containing syringes (BD vacutainer, CPTTM cell preparation tube, sodium citrate gel and density gradient media, Becton Dickinson, Heidelberg). Lymphocytes were isolated by centrifugation. For the analysis, 10^6 cells were lyzed on a membrane filter (25 mm diameter) and incubated for 50 min at 37°C with Fpg protein

(1 µg/ml) or buffer alone immediately before elution from the filter. The numbers of modifications sensitive to the repair glycosylase Fpg were obtained by subtraction of the number of single-strand breaks (SSB) observed in experiments without glycosylase treatment. Elution curves obtained with γ -irradiated cells were used for calibration, assuming that 6 Gy generated 1 single-strand break per 10⁶ bp.

Oxidative Status

As a marker for oxidative stress, the level of lipid peroxides (PerOx, Immundiagnostic, Bensheim, Germany) was determined. Lipid peroxides are assumed to result from reactions of lipids with free radicals. The test is based on the reaction of horseradish peroxidase with serum peroxides using tetramethylbenzidine (TMB) as a chromogen substrate (450 nm wavelength). In brief, 2.5 ml blood was coagulated for 30 minutes at room temperature, centrifuged at 3000 upm for 10-15 minutes and serum was frozen at -20°C. 10 µl of subject's serum and of calibrators were pipetted into a 96 well microplate and 100 µl reaction buffer was added. Optical density of the samples was measured at 450 nm in an ELISA reader. 100 µl reaction buffer mixture was added, followed by incubation at 37°C. Afterwards, 50 µl of stop solution was added. Subsequently, the second measurement of optical density of the sample was performed at 450 nm. The difference between first and second measurement was proportional to the peroxide content of the sample.

Antioxidative Enzymes

Superoxide Dismutase (SOD)

Superoxide dismutase (SOD) activity was measured photometrically using the colorimetric assay kit of Randox laboratories (Krefeld, Germany, kit no. SOD125). The principle of the test is based on the method of Woolliams et al. [36]. 500 µl whole blood supplemented with EDTA was used, from which cells were isolated by centrifugation and lyzed by a hypotonic buffer. The quantification method uses xanthine and xanthine oxidase to generate superoxide radicals. The superoxide anions react with 2-(4-iodophenyl)-3-(4nitrophenol)-5-phenyltretazolium chloride (INT) to form a red formazan dye. The SOD in the donor's blood sample competes with INT for superoxide radicals and thereby inhibits the production of the formazan dye. The degree of inhibition corresponds to the activity of SOD. Mean precision is 858.2 U/g Hb with a coefficient of variation (CV) of 12.2%.

Glutathione Peroxidase (GPx)

Glutathione peroxidase (GPx) activity was measured by photometry using the colorimetric assay kit of Randox laboratory (Krefeld, Germany, kit no. RS505). 50 μ l EDTA whole blood was used, in which cells were lyzed by addition of a hypotonic buffer. The principle of the test is based on the method of Paglia and Valentine [37]. GSH peroxidase catalyzes the oxidation of GSH by cumene hydroperoxide. In the presence of GSH reductase and NADPH, the oxidized GSH is converted to the reduced form accompanied by the oxidation of NADPH to NADP. At 340 nm wavelength, the decrease in absorbance is measured. The activity of GPx is assessed from the decrease in absorption at 340 nm wavelength due to oxidation of NADPH to NADP⁺. Mean precision is 43.9 U/g Hb; the CV is 5.6%.

<u>Total Glutathione</u>

Total glutathione (reduced and oxidized glutathione, GSH and GSSG) was measured photometrically by the method of the Medizinisches Labor (Bremen, Germany) as described in Richie *et al.* [38]. The principle is that reduced glutathione is oxidized to GSSG, which reacts with 5,5'-dithio-bis-(2-nitrobenzoe acid). Briefly, 2.7 ml frozen EDTA whole blood was used and samples were deproteinized by the addition of metaphosphoric acid (MPA). After protein precipitation, 5,5'-dithiobis-(2-nitrobenzoic acid) and glutathione oxidoreductase solution were added. The reaction was started by adding NADPH solution and the rate of color change was measured at 410 nm wavelength. Mean precision (n = 9) is 243.7 mg/L; coefficient of variation is 2.6%.

Antioxidative Micronutrients

<u>Vitamin C</u>

Vitamin C (µmol/L) was determined in plasma using the reagent kit-no. 65000 of Chromsystems (München, Germany) for HPLC analysis. Interfering components were removed by means of protein precipitation. For accurate quantification, a stable internal standard was used. In brief, the precipitation reagent was reconstituted with 1.5 ml internal standard. 100 µl of the precipitation reagent was pipetted into a reaction vial. 100 µl plasma was added. After incubation for 10 min at 4 °C and centrifugation for 5 min at 13000 rpm, 20 µl of the supernatant was injected into the HPLC system. Intra-assay CV was < 3% and inter-assay CV < 5%.

<u>Beta Carotene</u>

Beta carotene (µmol/L) was determined in serum using the reagent kit-no. 32000 of Chromsystems (München, Germany) for HPLC analysis. The sample was cleaned up by fast precipitation and extraction steps. In brief, 2.5 ml blood was coagulated for 30 minutes at room temperature, centrifuged at 3000 rpm for 10-15 minutes and aluminium foil was wrapped around. 100 µl serum was pipetted into a light protected reaction vial. 50 µl internal standard was added and mixed. 50 µl precipitation reagent was added and afterwards 200 µl extraction buffer. After centrifugation for 10 min at 13000 rpm, the supernatant was transferred into a light protected autosampler vial and 50 µl was injected into the HPLC system. Beta carotene was measured after extraction and HPLC separation at 460 nm wavelength. The analyte was quantified by the inclusion of an internal standard, which is a non-natural occurring carotenoid-derivative, so that a single detection wavelength is required. Intra- and inter-assay CV is < 5%.

<u>Zinc</u>

Zinc (μ mol/L) was measured by atomic absorption spectrometry (AAS). 2.5 ml blood was coagulated for 30 min at room temperature, centrifuged at 3000 rpm for 10-15 min. 500 μ l serum was deproteinized with TCA and was subjected to atomic absorption spectrophotometry. The concentration of zinc in the sample was computed from the calibration curves. Intra- and inter-assay CV is < 5 %.

Analysis of Nutritional and Life Style-Related Parameters

The assessment of nutritional and life style-related parameters was considered for the past two month prior to collection of blood samples. Subjects completed a questionnaire with respect to their smoking habits, alcohol consumption, physical activity level, and nutritional habits. Smoking habits were coded as 1, 2 or 3 corresponding to non-smokers, past smokers, or current smokers, respectively. In the case of smokers, the number of cigarettes per week was noted. With respect to alcohol consumption, weekly consumption of wine, separately for red and white wine, was questioned (number of glasses). The physical activity level was determined considering the hours of activity per week. Moderate and medium physical activity level was defined as performing intense exercise up to seven hours per week. For those subjects, who took their dogs out for a walk at a low space, up to fourteen hours per week were defined as moderate to medium physical activity. Exhaustive physical activity was defined as performing eight and more hours of intense exercise per week.

Statistical Analysis

All statistical analyses were carried out using the SPSS/PC for Windows statistical software package release 11.0 (SPSS Inc., Chicago). Data were tested for normal distribution using the Kolmogorov-Smirnov test. Levene's test was applied to analyze homogeneity of variances. Data were given as mean and standard deviation (SD). Comparisons between age groups and between sexes were performed with t-tests for independent samples. Pearson correlation coefficients were calculated to analyze relationships between biochemical parameters. For all analyses, a two-tailed p < 0.05 was taken as statistically significant.

RESULTS

Reproducibility and Intraindividual Variability of Oxidative DNA Damage Measures

Prior to the analysis of intergroup differences in oxidative DNA damage, reproducibility and intraindividual variability of the determinations were assessed (Fig. 1). The reproducibility of the quantification of Fpg-sensitive DNA modifications per 10⁶ bp was tested by analying two independent blood samples that were taken immediately after one another from the donor. Results show that the second measurements varied between 77% and 118% of first measurement (mean \pm S.D., 101 \pm 16%; n = 5 subjects). The *intrain*dividual variability of Fpg-sensitive DNA modifications per 10⁶ bp was measured in seven subjects. Blood samples were taken at two different days (time interval ranging from 7 to 93 days, mean 44 days). The second measurement was between 76% and 238% of the first measurement (mean \pm S.D.: 147 \pm 67%). There was no relationship between the absolute difference between both measurements and the time interval between taking the blood samples.

Differences Between Younger and Older Subjects

The comparative analysis of healthy younger and older subjects showed a tendency towards higher levels of DNA single-strand breaks (SSB) in older subjects (SSB, p = 0.085) (Fig. 2). Also, a higher level of serum peroxides ("oxidative status") was observed in older men (Table 1), but findings did not reach statistical significance. With respect to the antioxidative enzymatic defense system, the SOD activity was slightly, but not statistically significantly, reduced in older subjects. The GPx activity was significantly increased in older males (p < 0.05), but not in older females, compared to younger subjects. The SOD/GPx ratio, indicative of the relative rates of the first and second steps of antioxidative enzyme defense, was lower in older men than younger ones (Fig. 3). Older men displayed higher GSH-GSSG activities than younger ones, whereas women showed no age group differences. Regarding exogenous non-enzymatic antioxidative substances, the plasma concentration of vitamin C was similar in older and younger subjects, whereas intake of β -carotene has a tendency (p = 0.090) to be higher in elderly females (Table 1). Also, serum concentrations of zinc were higher in older men and women than in younger ones, but group differences were statistically non-significant (Table 1).







Fig. (2). Steady-state levels of single strand breaks (SSB) and Fpgsensitive DNA modifications (Fpg-mod) in lymphocytes from 19 young (24-28 yrs) and 19 old human donors (62-79 yrs). Males and females are pooled, because sex differences are not statistically significant. Older subjects show a trendency towards higher levels of oxidative DNA damage (n.s.). Columns represent means from 19 subjects \pm standard deviations. SSB, young *versus* old, p = 0.085; Fpg-mod, young *versus* old, p = 0.246.



Fig. (1). Reproducibility (panel **A**) and intraindividual variability (**B**) of Fpg-sensitive DNA modifications (Fpg $mod/10^6$ bp). In each panel, the second measurement value is given as a percentage of the first measurement by subject.

Fig. (3). Ratios of superoxide dismutase (SOD) activity (U/g Hb) and glutathione peroxidase (GPx) activity (U/g Hb) in whole blood samples of old and young male and female subjects. Older men show a tendency towards lowered SOD/GPx ratio (young *versus* old subjects: men, p = 0.097; women, p = 0.664).

 Table 1.
 Oxidative DNA Damage, Oxidative Status, Levels of Antioxidative Enzymes, and Degree of Physical Activity in 40

 Healthy Younger and Older Adults

	Males						Females					
Character	Young (n = 10)		Old (n = 10)		. 9		Young (n = 10)		Old (n = 10)			
	Mean	SD	Mean	SD	t"	р	Mean	SD	Mean	SD	- t*	р
Oxidative DNA Damage												
SSB/10 ⁶ bp	0.044	0.018	0.075	0.044	-1.98	0.064	0.052	0.026	0.058	0.036	-0.42	0.682
Fpg mod/10 ⁶ bp	0.242	0.089	0.276	0.080	-0.87	0.394	0.256	0.058	0.273	0.056	-0.79	0.441
Oxidative Status												
Serum peroxides (µmol/L)	38.2	37.7	59.7	37.3	-1.28	0.216	390.0	325.3	131.1	111.5	2.38	0.036
Antioxidative Enzymes and Glutathione												
SOD (U/g Hb)	1134.7	253.4	1075.1	211.3	0.57	0.575	1111.6	245.3	1061.0	277.9	0.43	0.671
Total glutathione (mg/L)	350.7	64.2	370.4	43.6	-0.80	0.432	313.9	40.0	315.8	64.6	-0.08	9.938
GPx (U/g Hb)	29.9	9.5	40.2	12.2	-2.12	0.048	37.7	11.4	36.3	9.1	0.29	0.779
SOD/GPx ratio	44.5	25.3	29.1	11.4	1.75	0.097	33.4	15.7	30.7	10.8	0.44	0.664
Antioxidative Micronutrients												
Vitamine C (µmol/L)	11.5	3.3	11.8	3.9	-0.18	0.856	14.2	7.6	13.5	2.6	0.28	0.785
β-carotene (µmol/L)	433.0	368.4	586.3	574.0	-0.71	0.486	412.0	238.3	710.7	470.3	-1.79	0.090
Zinc (µmol/L)	89.3	7.6	86.8	16.5	0.44	0.668	72.1	12.0	87.5	12.7	-2.78	0.012
Level of Physical Activity (hours/week)	4.70	2.39	6.45	3.16	-1.40	0.180	2.92	2.40	5.73	3.92	-2.07	0.052

Abbreviations: Fpg mod/10⁶ bp; Fpg sensitive DNA modifications/ 10⁶ base pairs; GPx, Glutathione peroxidase; SD, standard deviation; SOD, superoxide dismutase; SSB/10⁶ bp, single strand breaks/ 10⁶ base pairs.

^a Two-tailed *t*-test for independent samples.

Relationships Among Biochemical Parameters

The oxidative status did not correlate with the numbers of SSB per 10^6 bp (r = 0.011, p = 0.951) or Fpg-sensitive modifications per 10^6 bp (r = -0.002, p = 0.990). The level of oxidative DNA damage inversely correlated with the SOD activities (Fig. 4), but showed no relationship with GSH-GSSG or GPx activities. The level of oxidative DNA damage (Fpgsensitive modifications per 10^6 bp) negatively correlated with the SOD/GPx ratio in both sexes, being statistically significant in men (r = -0.50, p = 0.031), but not in women (r = -0.17, p = 0.528) (Fig. 5). The oxidative status level negatively correlated with total glutathione (r = -0.32, p = 0.048), but displayed no statistically significant relationship with SOD or GPx activities. The glutathione levels and GPx activities were positively correlated with each other (r =0.39, p = 0.013). Both the oxidative status and oxidative DNA damage did not show any significant relationship with plasma levels of vitamin C, ß-carotene, or zinc.

Relationships Between Biochemical and Lifestyle Parameters

Neither physical activity level nor smoking behavior was associated with the oxidative status, oxidative DNA damage, or antioxidant enzyme activities. Subjects drinking at least one glass of red wine every week displayed lower numbers for the oxidative status than abstinent subjects (106.1 ± 107.3)

versus 214.2 \pm 298.8µmol/L; t = 1.46, p = 0.160, n = 40), but did not differ in their levels of oxidative DNA damage (0.26 \pm 0.086 *versus* 0.26 \pm 0.045 Fpg-sensitive modifications per 10⁶ bp; t = -0.19, p = 0.851, n = 36).



Fig. (4). Relationship between oxidative DNA damage (Fpgsensitive modifications/ 10^6 bp) and superoxide dismutase activity (SOD) in healthy subjects (r = -0.36, p = 0.031). Fpg-mod: Fpgsensitive modifications.



Fig. (5). Relationship between oxidative DNA damage (Fpgsensitive modifications/ 10^6 bp) and SOD/GPx ratio in healthy subjects (men: r = -0.50, p = 0.031; women: r = -0.17, p = 0.528). Fpgmod: Fpg-sensitive modifications.

DISCUSSION

We examined the hypothesis that during natural aging, the oxidative stress increases, resulting in higher levels of oxidative DNA damage in the nucleus. The comparative analysis of older and younger healthy adults showed in agreement with our hypothesis, a tendency for higher levels of SSB and Fpg-sensitive modifications, which include 8oxo-dG, in older subjects, particularly in males. This is to our knowledge, the first study which investigated 8-oxo-dG damage in healthy elderly people using alkaline elution. Our observation confirms those of Barnett and King [39], who noted higher levels of DNA damage in older subjects by analyzing single-stranded DNA. Likewise, Dušinská et al. [40] found increased DNA damage with age as well as increased repair activity of the 8-oxoguanine DNA glycosylase (OGG1). The latter observation is consistent with findings demonstrated by transcriptional profiling that aging in the human frontal cortex is associated with increased expression of OGG1 [41].

We further found that the amount of DNA damage is inversely and statistically significantly related with SOD activity. A similar relationship has been noted in the rat brain [42]. As expected, older men had higher oxidative stress levels than younger ones, as measured by serum peroxide levels (oxidative status). Likewise, other authors reported for older subjects, increased levels of oxidative stress, as assessed by measuring the malondialdehyde level (MDA), which is one end-product of the lipid peroxidation process [27, 30, 31].

The hypothesis that the antioxidative capacity is lowered in older subjects is supported by our data for SOD, but not for total glutathione and GPx. By contrast, GPx activity was increased in older men. This is in line with findings of other studies indicating that older subjects had lowered SOD activities [25, 27, 29, 43] and increased GPx activities [27, 28, 44]. In this context, transcriptional profiling of the human frontal cortex has recently shown an increased expression of GPx with aging [41]. However, some authors observed in elderly subjects unchanged [25, 29] or decreased GPx activities [26, 31]. A decrease of SOD with aging might be due to a lowered metabolic rate and oxygen consumption generating less ROS, and thereby a lower demand for SOD. Since decreased SOD activity could also reflect Zn deficiency [45], we tested the subjects for their Zn level. We observed no relationship between SOD activity and Zn level (p > 0.05).

A decrease of SOD and an increase of GPx with aging might result in an imbalance in the antioxidative enzymatic defense system (in particular elevated levels of superoxide radicals). There is an evidence that a biological optimum exists for the ratio between SOD and GPx+CAT activities, which might be more relevant than the absolute activities of the enzymes themselves [46]. It has further been postulated that an altered SOD-to-(GPx+CAT) ratio affects gene expression by affecting the binding and/or availability of transcription factors to DNA [46]. We found that the SOD/GPx ratio was lowered, but not statistically significant, in older males compared to younger ones. Interestingly, we observed that subjects with lowered SOD/GPx ratio had increased oxidative DNA damage, as indicated by the level of Fpg sensitive DNA modifications. In line with this finding, studies on murine aging showed that alterations in this ratio correlated with free racial mediated cell damage as measured by lipid peroxidation. In contrast, no increased lipid peroxidation was detected in those organs where the SOD/GPx ratio was maintained during aging [24]. Furthermore, patients with Down syndrome had an altered SOD/GPx ratio, which is supposed to play a role in premature aging of this group [46, 47].

The oxidative status and oxidative DNA damage were not significantly affected by the plasma levels of vitamin C, B-carotene, or zinc. This confirms other studies, which did not find a relationship between oxidative DNA damage and vitamin C or B-carotene [48, 49]. In the present study, the younger subjects were not significantly higher in antioxidant supplements than the aged group. Both groups did not differ significantly in their plasma levels of vitamin C, ß-carotene, and zinc, except for higher zinc levels in older women. In contrast, other authors reported decreased plasma levels of vitamin C with aging [50]. This can be explained by insufficient dietary intake of this vitamin by elderly persons, because the plasma levels of vitamin C are correlated with dietary intake [51]. With respect to ß-carotene levels, conflicting results have been published. Compared to healthy younger adults, institutionalized elderly displayed lowered levels of B-carotene [52]. Considering the age-related decrease of activities of antioxidant enzymes, a sufficient supply of dietary oxidants seems to be important in the aged. Regular supplementation with antioxidants using recommended doses might reduce oxidative damage, which promotes cardiovascular disease, cancer, and diabetes [53]. Supplementation with moderate levels of antioxidant vitamins also reduces exercise-induced oxidative damage [54].

Exercise status has a profound effect on oxidative stress damage. It has been shown that exhaustive exercise causes oxidative stress, whereas moderate exercise may act as an antioxidant [55]. Muscle-damaging resistance exercise influences antioxidant enzyme activity, i.e. alters SOD activity, elevates CAT activity, and declines GPx activity [56]. In elderly people, long-term physical activity preserves antioxidant capacity and limits oxidative damage accumulation [57]. In the present study, older and younger subjects showed moderate to medium physical activity, but not exhaustive activity. The older subjects of both sexes displayed a somewhat higher level of physical activity than younger ones, but age group differences were statistically nonsignificant. Thereby, the physical activity level of the subjects was not associated with altered oxidative stress level, oxidative DNA damage, or antioxidant enzyme activities.

Finally, some methodological aspects are worth mentioning. First, the observed large intraindividual variability of Fpg-sensitive modifications in some volunteers might be explained by the time interval between first and second measurement and by the season of data collection. The time interval between first and second measurement ranged from 7 to 93 days. Subjects with a larger time interval displayed a larger difference between both measurements. Also, subjects with the first measurement in spring/summer and the second measurement in winter time showed a larger intraindividual variability. However, the first measurements, which were used for group comparisons between age groups and sexes, were all taken in the same season (spring), thereby not influencing the results of the group comparisons. Second, the level of serum peroxidases was even ten times higher in young females than in young males and three times higher in young females than in old ones. This indicates highest oxidative status in young females. This might be attributed to the interaction between female sex hormones and the test system used for measuring the level of lipid peroxides as a marker of oxidative stress. We assume that female sex hormones, which are highest in young women, interact with the analytical test system. This assumption is supported by our observation that pregnant women (not included in the present study) showed a higher level of oxidative status than non-pregnant women. Further studies are needed, which should analyze the level of serum peroxidases in a sample of younger women before, during, and after pregnancy, using the present test system. Also, the test results should be compared with those of a second test system. The test system for oxidative status might be one limitation for the present study. Also, the sample size of the present study is small. Further studies with more volunteers are needed.

CONCLUSION

Basic insights in genetic and cellular causes of natural aging may help to influence aging processes in a positive manner. The results of the present study give evidence for a tendency towards increased DNA damage (single strand breaks) in older males. Also, oxidative stress was elevated in older males, but results did not reach statistical significance. In addition, results indicate age-related changes in the balance between the first step (SOD) and second step (GPx) of the enzymatic antioxidant defense system. The SOD/GPx ratio was found to be related to the level of oxidative DNA damage.

ACKNOWLEDGEMENTS

We are very grateful to Mrs. Reghina Coman, MD, Institute of Sports Medicine, Gutenberg University of Mainz, for taking blood samples of the volunteers. We thank all volunteers very much for taking part in this study. The study was financially supported by a Forschungsfonds grant from the Gutenberg University of Mainz.

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Received: August 8, 2008

Revised: October 22, 2008

Accepted: October 29, 2008

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