

## A preliminary assessment of singlet oxygen scavenging, cytotoxic and genotoxic properties of *Geranium macrorrhizum* extracts\*

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**Strong radical-scavenging activity of *Geranium macrorrhizum* extracts isolated by using various solvent systems has been reported previously. This study aimed at expanding the knowledge on the bioactivities of antioxidatively active *G. macrorrhizum* butanol fraction, which was isolated from ethanolic extract (EB), and water fraction, which was isolated from water extract (WW) by measuring their singlet oxygen scavenging properties, as well as preliminary assessment of cytotoxicity and genotoxicity toward mammalian cells. The cytotoxicity (necrosis induction) of the extracts in bovine leukemia virus-transformed lamb kidney fibroblasts (line FLK) was partly prevented by antioxidants and stimulated by the prooxidant BCNU (*N,N'*-bis(2-chloroethyl)-*N*-nitrosourea). This indicates that the cytotoxicity of *G. macrorrhizum* extracts is at least partly attributed to their prooxidant action, presumably due to the formation of quinoidal products of their (auto)oxidation. The latter was evidenced by the nature of the peroxidase-catalyzed oxidation products, which supported DT-diaphorase-catalyzed oxidation of NADPH and participated in conjugation reactions with reduced glutathione. The genotoxic properties were studied using chromosome aberration (CA) and sister chromatid exchange (SCE) tests in human lymphocytes *in vitro* and *Drosophila melanogaster* somatic mutation and recombination test (SMART) *in vivo*. In the CA test, only the highest doses of both fractions significantly increased chromosome aberration frequency. In the SCE test, both fractions induced SCEs in a clear dose-dependent manner. *G. macrorrhizum* extracts were not genotoxic in the SMART test *in vivo*. Our data indicate that in spite of the possible beneficial (antioxidant) effects of *Geranium* extracts, the possibilities of their use as ingredients of functional foods and/or food supplements should be further examined due to their cytotoxic and genotoxic effects resulting mainly from the action of quercetin-derived components abundant in the extracts.**

**Keywords:** *Geranium macrorrhizum*, polyphenols, cytotoxicity, genotoxicity

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### INTRODUCTION

Aromatic and medicinal plants are a good source of bioactive compounds which may be valuable ingredients for various applications, including pharmaceuticals, func-

tional foods and food supplements. With the advent of modern analytical and bioactivity assessment techniques studies of plant-derived natural preparations and purified compounds has expanded substantially; however, considering the vast biodiversity of the plant kingdom and the numerous factors affecting the production of secondary metabolites in plants this area of research is still not sufficiently explored. Thus, our previous studies of aromatic and medicinal plants grown in Lithuania by using various methods resulted in the evaluation of antioxidative properties of some less investigated plants (Dapkevicius *et al.*, 1998; Weel *et al.*, 1999; Bandonienė *et al.*, 2000; Povilaitytė & Venskutonis, 2000) and identification of new natural antioxidants (Dapkevicius *et al.*, 2002; Pukalskas *et al.*, 2002; Miliuskas *et al.*, 2004a; 2005). Extracts of *Geranium macrorrhizum* were shown to possess powerful radical scavenging capacity (Miliuskas *et al.*, 2004b) and antioxidant activity, which were due to the presence of the identified polyphenolic components (Miliuskas *et al.*, 2004c). The antioxidative properties of *G. macrorrhizum* extracts added as natural antioxidants were further evaluated in Dutch-style fermented sausages (Miliuskas *et al.*, 2007). *G. macrorrhizum* is a grassy perennial plant with long roots, which is used for the production of essential oil containing mainly the sesquiterpene garmacrone. The plant is known to be rich in tannins and its extracts have been reported to possess a broad spectrum of antimicrobial activities. In addition it was declared to have strong hypotensive, astringent activity, as well as cardiogenic and sedative properties (Bates-Smith, 1981; Ivancheva *et al.*, 1992).

The bioactivities of plant extracts are partially associated with the presence of various polyphenolic compounds, which may scavenge free radicals, thus acting as primary antioxidants as well as quenching and reacting with singlet oxygen (<sup>1</sup>O<sub>2</sub>) (Tournaire *et al.*, 1993; Darmann *et al.*, 1998; Bensasson *et al.*, 1999). However, apart from their beneficial properties, polyphenols may be tox-

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**Abbreviations:** AlPcS<sub>4</sub>, Al-phthalocyanine tetrasulphonate; BCNU, *N,N'*-bis(2-chloroethyl)-*N*-nitrosourea; cI<sub>50</sub>, compound concentration doubling the photohemolysis time of erythrocytes; cL<sub>50</sub>, compound concentration causing 50% cell death; CA, chromosome aberration; DMSO, dimethyl sulfoxide; EB, ethanol/butanol fraction; NQO1, NAD(P)H:quinone oxidoreductase; POD, peroxidase; RI, refractive index; ROS, reactive oxygen species; SCE, sister chromatid exchange; SMART, somatic mutation and recombination test; WW, water/water fraction.

ic in mammalian cells among other modes of their action by exerting oxidative stress-type cytotoxicity. This is attributed to the polyphenol autooxidation in cell growth media with the production of extracellular  $H_2O_2$  (Nakagawa *et al.*, 2004, and references therein), the intracellular generation of reactive oxygen species (Galati & O'Brien, 2004, and references therein), and the depletion of intracellular reduced glutathione (GSH) and other reduced thiols by the quinone/quinomethide-type (auto)oxidation products of polyphenols (Boersma *et al.*, 2000).

Plant extracts are very complex mixtures often containing toxic as well as antimutagenic compounds. Therefore, studies on their "latent" toxicities such as teratogenesis, carcinogenesis and mutagenesis are of particular relevance. The genotoxic effects of plant extracts have been studied over the years and it was shown that a great number of plants that are used as food ingredients or in traditional medicine possess mutagenic properties (Lazutka *et al.*, 2001; Cardoso *et al.*, 2006; Deciga-Campos *et al.*, 2007; Verschaevé & van Staden, 2008). Considering the growing use of plant extracts, genotoxicity testing and identification of genotoxic compounds could help to increase their safety.

The objectives of the present study were to evaluate the singlet oxygen scavenging, cytotoxicity and genotoxic activities of selected fractions of *G. macrorrhizum* extracts. For the latter purpose we used chromosome aberration and sister chromatid exchange tests in human lymphocytes *in vitro* and *Drosophila melanogaster* wing spot somatic mutation and recombination test (SMART) *in vivo*.

## MATERIALS AND METHODS

**Preparation of extracts and their fractions.** Dried plant material of *G. macrorrhizum* was obtained from the Kaunas Botanical Garden of Vytautas Magnus University (Lithuania). The extracts and their fractions were prepared from 100 g of dry ground mass by step-wise extraction as described previously (Miliauskas *et al.*, 2004c). Briefly, herb was extracted with  $2 \times 500$  ml of *tert*-butyl methyl ether (TBME) at room temperature for  $2 \times 12$  h under constant stirring with a magnetic stirrer. The TBME extract was filtered through a medium porosity filter paper and the solvent was evaporated in a rotary evaporator. The residue of plant material remaining after extraction with TBME was re-extracted under the same conditions with ethanol (96%, v/v) and finally with  $H_2O$ . The extracts obtained with EtOH and  $H_2O$  were further partitioned between water and 1-butanol resulting in ethanol/butanol (EB), ethanol/water (EW), water/butanol (WB) and water/water (WW) fractions. The EW and WW fractions were freeze-dried. Seven phenolic compounds, namely gallic acid, ellagic acid, 4-galloyl quinic acid, the flavonoid quercetin and three of its glycosides, quercetin-3- $\beta$ -glucopyranoside, quercetin-3- $\beta$ -galactopyranoside and quercetin-4'- $\beta$ -glucopyranoside were isolated and identified in the various fractions (Miliauskas *et al.*, 2004c).

**Enzymatic reactions.** Rat liver NAD(P)H:quinone oxidoreductase (DT-diaphorase, NQO1, EC 1.6.99.2) was prepared as described elsewhere (Prochaska, 1988), while horseradish peroxidase (POD, EC 1.11.1.7) was obtained from Sigma. The enzyme concentrations were determined spectrophotometrically according to  $\epsilon_{460} = 11 \text{ mM}^{-1} \cdot \text{cm}^{-1}$  (NQO1), and  $\epsilon_{402} = 102 \text{ mM}^{-1} \cdot \text{cm}^{-1}$  (POD). All other reagents were obtained from Sigma and used as received. Enzymatic reactions carried out in 0.1 M

K-phosphate (pH 7.0) with 1 mM EDTA at 25°C were studied spectrophotometrically using a Hitachi-557 UV-VIS spectrophotometer. To characterize the spectral changes during the POD-catalyzed oxidation, the spectra of 10–50  $\mu\text{g/ml}$  of extracts were recorded at 250–700 nm before and after the incubation with 0.1  $\mu\text{M}$  POD and 100–300  $\mu\text{M}$   $H_2O_2$  for 5 min. In separate experiments, 5.0 mM GSH was added at the end of the reaction. The activity of the extract oxidation product(s) in supporting NQO1-catalyzed NADPH oxidation was studied as follows: after the oxidation of extracts by POD +  $H_2O_2$ , the excess  $H_2O_2$  was decomposed by 50 U/mg catalase. Subsequently, 100  $\mu\text{M}$  NADPH and 20 nM of NQO1 were introduced into the reaction mixture, and the NADPH oxidation rate was determined according to  $\Delta\epsilon_{340} = 6.2 \text{ mM}^{-1} \cdot \text{cm}^{-1}$ . In control samples in the absence of NQO1 the oxidation of NADPH by the extract oxidation product(s) did not take place.

**Mammalian cell cytotoxicity studies.** Bovine leukemia virus-transformed lamb kidney fibroblasts (line FLK) were grown and maintained in Eagle's medium supplemented with 10% fetal bovine serum at 37°C as described previously (Nemeikaitė-Čėnienė *et al.*, 2005, and references cited therein). Cells ( $2.5 \times 10^4/\text{ml}$ , final volume, 5 ml) were seeded on 18 mm  $\times$  18 mm glass slides in 5 ml flasks in the presence or in the absence of extracts and additional compounds and were grown for 24 h. The extracts and additional compounds were dissolved in DMSO, and 5–10  $\mu\text{l}$  of DMSO solution was added into the cell growth medium. In control experiments, this amount of DMSO did not affect cell viability. Further, the slides were rinsed with phosphate-buffered saline, and stained with Trypan Blue. The viable and nonviable (necrotic Trypan Blue-accumulating) cells on the slides were counted under a light microscope.

**Erythrocyte photohemolysis protection studies.** Freshly prepared suspension of human erythrocytes (Vilnius Blood Transfusion Centre) was stored in 0.01 M K-phosphate with 0.137 M NaCl, 2.7 mM KCl, 10 mM glucose, and 1 mM EDTA as described previously (Marozienė *et al.*, 2000). For photolysis experiments, erythrocytes were diluted with the same solution to  $2.5\text{--}2.6 \times 10^6/\text{ml}$ . Afterwards, 10  $\mu\text{M}$  Al-phthalocyanine tetrasulphonate (Al-PcS<sub>4</sub>, Porphyrin Products, Logan, UT, USA) and extract solution were added to a stirred suspension of erythrocytes in the dark, 30 min prior to irradiation. The irradiation was performed at 25°C under continuous stirring, in a 1.0 cm inner diameter glass cell. The light source was a 250 W tungsten lamp with a cut-off filter ( $\lambda > 590$  nm) and a fluence rate of 25 W/m<sup>2</sup>. During continuous irradiation aliquots of erythrocytes (0.1 ml) were taken and diluted 20 times. The degree of their lysis was measured according to the decrease in their apparent absorbance at 740 nm caused by cell disruption (Sorata *et al.*, 1988). Typically, the erythrocyte lysis was characterized by a lag-phase of 40–60 min, which was defined as the intercept of the tangent to the turning point of the lysis curve with the time axis. The efficiency of compounds in the protection of erythrocytes from photohemolysis was expressed as the concentration giving a 2-fold increase in the lag-phase ( $CI_{50}$ ). It was calculated from lag-phase *vs.* extract concentration plots. Complete hemolysis was obtained by addition of 40  $\mu\text{g/ml}$  digitonin to erythrocytes.

**Cytogenetic test in human lymphocytes *in vitro*.** Whole peripheral blood from healthy volunteer was incubated at 37°C for 72 h in Hepes-buffered RPMI 1640 medium supplemented with 12% heat-inactivated new-

born calf serum, 7.8 µg/ml phytohemagglutinin P, 50 µg/ml gentamycin, 10 µg/ml 5-bromo-2'-deoxyuridine. All reagents used for the cell culture were purchased from Sigma. Treatment with plant extracts was carried out 48 h after culture initiation and lasted for 24 h. As whole blood cultures display many properties common with the liver microsomal cytochrome P450 system (Starke & Mieczal, 1989), no external metabolising enzymes were added. Test solutions were prepared in RPMI 1640 medium (WW) or dissolved in ethanol (EB) and then diluted with RPMI 1640 medium to the desired concentration. Working solutions of extracts were made just before treatment. Two parallel cultures were used for each concentration of plant extracts tested. Ethanol at a final concentration of 7.5 µl/ml was used as a solvent control for the EB fraction. Ethanol concentration in the experimental series did not exceed this concentration. Two cultures were left untreated and served as a blank control. The cultures were exposed to colchicine at a final concentration of 0.6 µg/ml for the last 3 h of incubation. The cells were harvested, hypotonically swollen in 0.075 M KCl and fixed in methanol/acetic acid (3:1, v/v). Air-dried slides were differentially stained by fluorescence plus Giemsa technique as described previously (Lazutka, 1991). Briefly, the slides were stained for 10 min with 10 µg/ml of Hoechst 33258 dye (dissolved in 0.07 M Soerensen's buffer, pH 6.8). Then the slides were rinsed, wetted with citrate buffer (pH 8.5), covered with cover slips and exposed to UV light (400 W mercury lamp at a distance of 15 cm) for 6–7 min. The slides were then rinsed and stained for 3–4 min with 5% Giemsa. Cytogenetic analysis was performed on coded slides. No less than 100 first-mitotic division metaphases per culture were analysed for chromosome aberrations, and no less than 50 second-division metaphases for sister chromatid exchanges. Aberrations were scored as individual types, but for convenience they were grouped as chromatid breaks (ctb), chromatid exchanges (cte), chromosome breaks (csb) and chromosome exchanges (cse).

Cell replicative kinetics was determined by means of replicative index (RI)=[M<sub>1</sub> + 2M<sub>2</sub> + 3M<sub>3</sub>]/N, where M<sub>1</sub>, M<sub>2</sub>, M<sub>3</sub> are the numbers of cells that had undergone one, two or three cycles of replication, and N is the total number of cells scored). Three hundred cells were scored to determine RI.

**Somatic mutation and recombination test (SMART) in *Drosophila melanogaster* in vivo.** The basic procedures for the *D. melanogaster* somatic mutation and recombination test (SMART) were performed according to Graf *et al.* (1984) and Graf and van Schaik (1992). Offspring from a cross of ORR/ORR; *flr<sup>3</sup>/In(3LR) TM3, ri<sup>p</sup> sep l(3)89Aa bx<sup>34c</sup> e Bd<sup>8</sup>* females with *mvb/mvb* males were used for mutation analysis (both strains kindly provided by Dr H. Frei, Zurich, Switzerland). The ORR strain has chromosomes 1 and 2 from a DDT-resistant Oregon strain characterized by constitutively over-expressed *CYP450* genes. The *CYP450* enzymes play the main role in the bioactivation of xenobiotics, as well as natural products including phytochemicals (van Iersel *et al.*, 1999). Thus, the use of this high-bioactivation strain of *D. melanogaster* facilitates the detection of promutagens of numerous classes of compounds. The markers *mvb* (multiple wing hairs) and *flr<sup>3</sup>* (misshapen, flare-like hairs) are recessive wing-hair mutations located on the third chromosome at genetic map positions 0.3 and 38.8, respectively.

Eggs from the crosses were collected over 10 h periods. Progeny were raised on Instant *Drosophila* Medium

(No. D7670, Sigma) at 25 °C. Two schedules of larvae treatment were applied. Firstly, the crossed flies were permitted to lay eggs for 10 h in vials containing medium prepared with the test solution. Thus, the exposure duration was about 120 h (i.e., the period from fly egg till case-worm). According to the second schedule, the 72 ± 5 h-old larvae were exposed to plant extracts by adding test solution to the surface of the medium and were fed on this medium for the rest of their development, during approx. 48 h. The test solutions were prepared in distilled water (WW fraction) or dissolved in ethanol (EB fraction) and then diluted with distilled water to the desired concentration, with the final ethanol concentration 5% (v/v). The solutions were always prepared immediately before use. Solvent controls were included in all treatments. Trans-heterozygous (*mvb flr<sup>3</sup>/mvb<sup>+</sup> flr<sup>3</sup>*) flies were collected and stored in 70% ethanol. The wings of adult flies were mounted on glass slides in Faure's solution, coded and scored under a magnification of 400× (Nikon E200) for the presence of single (*mvb* or *flr<sup>3</sup>* phenotype) or twin (adjacent *mvb* and *flr<sup>3</sup>* clones) spots. The spots were recorded according to standard procedures (Graf *et al.*, 1984). No less than 40 wings were analysed per each experimental point.

**Statistical analysis.** Statistical analyses were performed using InStat V2.02 (GraphPad Software, CA, USA) statistical package. Statistical tests were chosen according to the nature of the data analysed. X<sup>2</sup>-test with Yate's correction was used to estimate the induction of chromosome aberrations and somatic mutations. A one-way analysis of variance (ANOVA) and the Student's two-sided *t*-test were used for the evaluation of SCE occurrence, and  $\chi$ -test for RI analysis. *P* < 0.05 was considered as the level of significance.

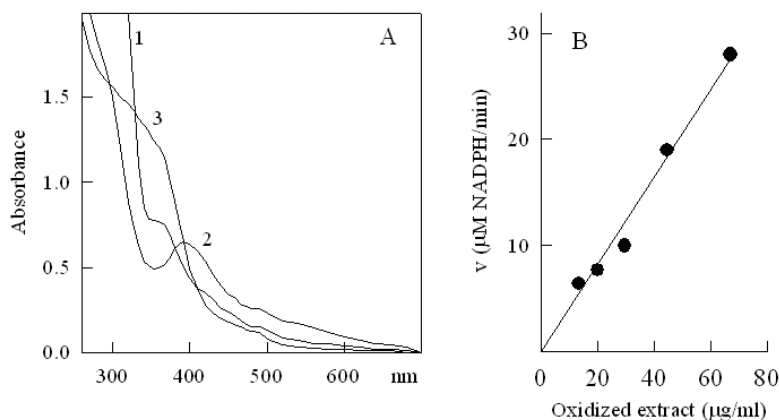
## RESULTS AND DISCUSSION

### Protective effects of *G. macrorrhizum* extracts against <sup>1</sup>O<sub>2</sub>-mediated photohemolysis of erythrocytes

In order to extend previously reported data on the antioxidant properties of *G. macrorrhizum* extracts (Miliauskas *et al.*, 2004c), we analyzed their capacity to neutralize the damaging action of singlet oxygen (<sup>1</sup>O<sub>2</sub>), using human erythrocyte photosensitized lysis by Al-phthalocyanine tetrasulphonate (Al-PcS<sub>4</sub>) (Marozienė *et al.*, 2000). Earlier studies showing the absence of photohemolysis protection by catalase and superoxide dismutase, and protective effects of azide, a specific quencher of <sup>1</sup>O<sub>2</sub>, have demonstrated that <sup>1</sup>O<sub>2</sub> plays a major role in the erythrocyte lysis (Marozienė *et al.*, 2000). For *G. macrorrhizum* extracts, the cI<sub>50</sub> values calculated from the data in Table 1 were equal to ≥300 µg/ml (EB fraction) and ≥380 µg/ml (WW fraction). Among the polyphenols

**Table 1. Protective effects of *G. macrorrhizum* EB and WW fractions against Al-PcS<sub>4</sub>-induced photohemolysis of erythrocytes** *n* = 3, *P* < 0.02 for (2–4) vs. (1).

No.	Additions (µg/ml)	lag-period (min)	
		a) EB	b) WW
1.	None	45 ± 6.0	51 ± 7.0
2.	100	60 ± 8.0	62 ± 5.0
3.	200	72 ± 6.0	74 ± 6.0
4.	300	88 ± 8.0	89 ± 7.0



**Figure 1.** The activity of *G. macrorrhizum* extracts towards redox enzymes. (A) Absorbance of extract (50 μg/ml) before (1) and after addition of 1.0 μM peroxidase and 100 μM H<sub>2</sub>O<sub>2</sub> (2), and after subsequent addition of 5 mM GSH (3); (B) oxidation of NADPH (100 μM) by 0.02 μM flavoenzyme DT-diaphorase (NAD(P)H:quinone oxidoreductase).

identified in *G. macrorrhizum* extracts, quercetin possessed a  $cI_{50} = 170 \pm 30 \mu\text{M}$ , which corresponds to  $51 \pm 9.0 \mu\text{g/ml}$ , whereas gallic acid was a much less effective protecting agent ( $cI_{50} \geq 2.0 \text{ mM}$ , or  $\geq 340 \mu\text{g/ml}$  (Marozienė *et al.*, 2000)). Thus, one may expect that quercetin but not gallic acid plays a major role in the protective effects of *G. macrorrhizum* extracts against <sup>1</sup>O<sub>2</sub>. This also may be valid for their glycoside derivatives, because the presence of a carbohydrate moiety decreases the lipophilicity of polyphenolic compounds (Nemeikaitė-Čėnienė *et al.*, 2005) and in turn decreases their protection efficiency against the photosensitized erythrocyte lysis (Marozienė *et al.*, 2000).

#### Prooxidant mammalian cell cytotoxicity of *G. macrorrhizum* extracts

As one of the cytotoxicity mechanisms of polyphenols is associated with the formation of their prooxidant (auto)oxidation products of quinoidal structure, first we examined the possibility of formation of quinoidal oxidation products in *G. macrorrhizum* extracts. The oxidation of *G. macrorrhizum* EB fraction by a model system, H<sub>2</sub>O<sub>2</sub> and peroxidase (POD) (Metodiewa *et al.*, 1999), leads to a disappearance of the absorbance shoulder at 360 nm and to the formation of a new absorbance peak at 400 nm (Fig. 1A). The subsequent addition of GSH leads to the formation of new product(s) possessing an absorbance shoulder at 350 nm (Fig. 1A). This shows that polyphenolic components of *G. macrorrhizum* extracts may be oxidized into quinone-type compounds which may react with GSH forming GS-substituted hydroquinones (Boersma *et al.*, 2000; Galati & O'Brien, 2004). The formation of quinone-type compounds during the POD-catalyzed oxidation of polyphenolic components of *G. macrorrhizum* extracts is confirmed by the dose-dependent effect of the oxidised extract on DT-diaphorase (NAD(P)H:quinone oxidoreductase, NQO1, EC 1.6.99.1)-catalyzed NADPH oxidation (Fig. 1B). This reaction is almost completely inhibited by 20 μM dicumarol, a specific inhibitor of NQO1. This definitely shows that quinoidal compounds may be formed during the (auto)oxidation of polyphenolic components of *G. macrorrhizum* extracts.

In the course of preliminary studies of the cytotoxicity (necrosis induction) by *G. macrorrhizum* extracts we used a bovine leukemia virus-transformed lamb embryo kidney fibroblasts culture (line FLK), which serves as a reference base in our investigation of prooxidant compounds (Nemeikaitė-Čėnienė *et al.*, 2005, and references therein). In this case, 98–99% of cells adhering to the glass sides did not accumulate Trypan Blue, i.e., they

were viable. The  $cI_{50}$  (compound concentration causing 50% cell death) was equal to  $112 \pm 15 \mu\text{g/ml}$  (EB fraction) or  $63 \pm 7.5 \mu\text{g/ml}$  (WW fraction). For comparison, in FLK cells the  $cI_{50}$  of quercetin was equal to  $42 \pm 5.4 \mu\text{g/ml}$  ( $140 \pm 18 \mu\text{M}$ ), and that for gallic acid was equal to  $128 \pm 12 \mu\text{g/ml}$  ( $750 \pm 70 \mu\text{M}$ ) (Nemeikaitė-Čėnienė *et al.*, 2005). Again, this shows that the cytotoxicity of *G. macrorrhizum* extracts may be mainly attributed to the action of quercetin and its derivatives, but not to those of gallic acid. The prooxidant nature of the cytotoxicity of *G. macrorrhizum* EB fraction is further confirmed by the data in Table 2, which show that the extract cytotoxicity was prevented by the antioxidant *N,N'*-diphenyl-*p*-phenylene diamine (DPPD) and by the Fe-ion chelator desferrioxamine, the latter preventing the Fenton reaction, but was stimulated by BCNU (*N,N'*-bis(2-chloroethyl)-*N*-nitrosourea) which acts as a prooxidant through inactivating the antioxidant enzyme glutathione reductase (EC 1.6.4.3) (Ollinger & Brunmark, 1991; Nemeikaitė-Čėnienė *et al.*, 2005). Considering other modes of the cytotoxicity of polyphenolic plant extracts, it is worth noting that the induction of apoptosis, a programmed cell death, in several cell lines by the isolated polyhydroxybenzenes and flavonoids was caused by ROS formation, also being partly prevented by DPPD and desferrioxamine (Nemeikaitė-Čėnienė *et al.*, 2009, and references therein). This also points to the prooxidant character of their apoptosis induction. However, further studies should be performed in this direction taking into account parallel apoptosis induction mechanisms of polyphenols, which apparently do not involve the formation of reactive oxygen species (Hsu *et al.*, 2007; Chien *et al.*, 2009).

#### Genotoxicity of *G. macrorrhizum* extracts

Genotoxicity in human lymphocytes *in vitro* was assessed using two endpoints — chromosome aberrations and sister chromatid exchanges. Analysis of chromosome aberrations is widely accepted as a test for clastogenicity. Although the exact mechanism that leads to an increased exchange of segments between sister chromatids is still not known, sister chromatid exchanges are successfully adopted as a test for genotoxicity. Both tests are used in genotoxicity assessment of plant-derived chemicals (Stopper *et al.*, 2005). The EB as well as WW fractions of *G. macrorrhizum* extract induced sister chromatid exchanges in a clear dose-dependent manner (Table 3). The dose-response relationships may be described by linear equations:  $Y = 8.24 + 0.08X$  ( $r^2 = 0.96$ ) for EB fraction and  $Y = 10.09160 + 0.07X$  ( $r^2 = 0.98$ ) for the WW fraction, where Y is the number of sister chromatid exchanges per cell, X is the concentration of the extract (μg/ml). Treat-

**Table 2. The cytotoxicity of *G. macrorrhizum* EB extract.** Effects of antioxidant *N,N'*-diphenyl-*p*-phenylene diamine (DPPD), Fe-ion chelator desferrioxamine, and potentiating effect of prooxidant alkylating agent *N,N'*-bis (2-chloroethyl)-*N*-nitrosourea (BCNU)

Additives	Cell viability (%) ( <i>n</i> = 3–4, <i>P</i> < 0.02)
<i>G. macrorrhizum</i> EB, 110 µg/ml	49.2 ± 2.5
Extract + DPPD (3.0 µM)	70.3 ± 3.5
Extract + desferrioxamine (300 µM)	72.8 ± 4.9
Extract + BCNU (20 µM)	29.2 ± 3.9

DMSO, 5–10 µL in 5 ml growth media (negative control) decreased cell viability by < 1%.

ment with extracts did not inhibit cell replicative kinetics. In the chromosome aberration test, the WW fraction caused a significant increase of chromosome aberrations at the highest concentration (200 µg/ml) only (Table 4). The EB fraction of *G. macrorrhizum* extract was the most clastogenic at a concentration of 150 µg/ml (more than eightfold increase over ethanol control), while at a higher concentration the number of aberrant cells decreased (Table 4). The prevalent type of induced aberrations in the case of both WW and EB fractions was chromatid

breaks. Interestingly, that genotoxicity profile of the EB fraction in human lymphocytes *in vitro* was very similar to that one reported for another plant-derived complex mixture, essential oil from peppermint herb (Lazutka *et al.*, 2001).

Pooled data of the wing spot analysis are reported in Table 5. It should be noted that there were very few spots, predominantly single small ones (not shown). Water controls showed background frequencies of 16.3–20.4% wings with spots and 0.18–0.20 total spots per wing. These values are within the range previously reported in the literature (0.16 to 0.60 total spots per fly) (Tellez *et al.*, 2007; Castro *et al.*, 2008). The frequencies of wings with spots and total spots in ethanol controls did not significantly differ from those in the water controls. No statistically significant difference between respective controls and different concentrations of the test solutions was found after 48 h or 120 h treatment. A statistically significant increase in the total number of spots was observed only among the flies treated with 1% WW fraction with exposure duration of 120 h, compared with the spot frequency in the water control.

The data presented in this study concern a genotoxic assessment of two fractions of *G. macrorrhizum* extract. The use of *in vitro* and *in vivo* assays was decisive in or-

**Table 3. Effects of different fractions of *G. macrorrhizum* extract on the frequency of sister chromatid exchanges (SCE) and replication index (RI) in human lymphocyte cultures**

Concentration of test solution (µg/ml)	Water/water (WW) fraction		Ethanol/butanol (EB) fraction	
	SCE/cell ± S.E.M.	RI ± S.E.M.	SCE/cell ± S.E.M.	RI ± S.E.M.
Blank	8.76 ± 0.49	2.47 ± 0.05		
Ethanol (7.5 µl/ml)			9.34 ± 0.41	2.15 ± 0.06
25	11.68 ± 0.65	2.31 ± 0.05	10.84 ± 0.55	2.59 ± 0.04
50	14.18 ± 0.62 <sup>a</sup>	2.44 ± 0.05	10.58 ± 0.49	2.33 ± 0.05
100	17.46 ± 1.02 <sup>a</sup>	2.36 ± 0.05	14.86 ± 0.79 <sup>a</sup>	2.37 ± 0.05
150	21.42 ± 1.00 <sup>a</sup>	2.36 ± 0.05	20.60 ± 1.19 <sup>a</sup>	2.27 ± 0.05
200	23.04 ± 0.88 <sup>a</sup>	2.46 ± 0.03	24.42 ± 1.08 <sup>a</sup>	2.23 ± 0.05

<sup>a</sup>*P* < 0.05 as compared to appropriate solvent control

**Table 4. Induction of chromosome aberrations in human lymphocytes by different fractions of *Geranium macrorrhizum* extract**

Treatment	Concentration (µg/ml)	Number of metaphases analysed	CA <sup>a</sup> per 100 cells (± S.E.M.)	Number of aberrations <sup>a</sup>			
				ctb	cte	csb	cse
Blank	0	100	1.0 ± 0.99	1	0	0	0
Ethanol	7.5	100	2.0 ± 1.4	2	0	0	0
Water/water (WW) fraction	25	100	3.0 ± 1.7	3	0	0	0
	50	100	4.0 ± 1.9	3	0	1	0
	100	100	1.0 ± 0.99	1	0	0	0
	150	100	2.0 ± 1.4	2	0	0	0
Ethanol/butanol (EB) fraction	200	100	8.0 ± 2.7 <sup>b</sup>	7	0	1	0
	25	100	7.0 ± 2.5	7	0	0	0
	50	100	6.0 ± 2.4	2	0	2	2
	100	100	4.0 ± 1.9	2	1	1	0
	150	100	17.0 ± 3.8 <sup>b</sup>	16	0	1	0
200	100	7.0 ± 2.5	7	0	0	0	

<sup>a</sup>CA, chromosome aberration; ctb, chromatid breaks; cte, chromatid exchanges; csb, chromosome breaks; cse, chromosome exchanges; <sup>b</sup>*P* < 0.05 as compared to appropriate solvent control.

**Table 5. Results of *Drosophila* wing spot test after treatment with different fractions of *G. macrorrhizum* extracts**

Concentration of test solution (%)	Water/water (WW) fraction				Ethanol/butanol (EB) fraction			
	Exposure duration 48 h		Exposure duration 120 h		Exposure duration 48 h		Exposure duration 120 h	
	Wings with spots (%±S.E.M.)	Spots per wing (m±S.E.M.)	Wings with spots (%±S.E.M.)	Spots per wing (m±S.E.M.)	Wings with spots (%±S.E.M.)	Spots per wing (m±S.E.M.)	Wings with spots (%±S.E.M.)	Spots per wing (m±S.E.M.)
Water	20.4±6.1	0.20±0.06	16.3±5.3	0.18±0.05	20.4±6.1	0.20±0.06	16.3±5.3	0.18±0.05
Ethanol (5%)	23.9±6.3	0.24±0.06	21.2±5.9	0.21±0.06	23.9±6.3	0.24±0.06	21.2±5.9	0.21±0.06
0.1	22.4±6.0	0.27±0.06	20.4±5.7	0.20±0.06	22.4±6.0	0.23±0.06	16.3±5.3	0.16±0.05
0.5	26.5±6.3	0.35±0.07	22.4±5.9	0.22±0.06	17.0±5.5	0.17±0.05	28.6±6.4	0.33±0.07
1	28.6±6.4	0.35±0.07	31.3±6.7	0.35±0.07 <sup>a</sup>	15.5±5.4	0.15±0.05	22.4±6.0	0.23±0.06

<sup>a</sup>*P*<0.05 as compared to appropriate control

der to cover different types of genotoxic damage. Only a slight (though significant at the highest concentrations) increase of chromosome aberrations was observed in human lymphocytes *in vitro*. However, both EB and WW fractions turned out to be potent inducers of SCEs. The dose-response relationships were linear for both fractions. The *G. macrorrhizum* extract is a complex mixture containing different phytochemical components. As it was mentioned above, gallic, ellagic and 4-galloyl quinic acids, quercetin and its glycosides have been identified in *G. macrorrhizum* extracts. These substances possess strong antioxidative activity (Miliauskas *et al.*, 2004c). However, previous studies have shown quercetin to have a dual effect. Low doses of quercetin have been shown to protect against oxidative injuries, while higher doses induced DNA damage, gene mutations and chromosome aberrations (Skibola & Smith, 2000; Stopper *et al.*, 2005). Our results suggest that the genotoxicity of *G. macrorrhizum* extract in human lymphocytes *in vitro* can be attributed to a pro-oxidant effect of quercetin. This proposition is substantiated by the findings of other researchers. The mutagenicity of four commercial ethanolic plant extracts (Tinctura Alchemillae, Extractum Crataegi, Extractum Myrtilli, Tinctura Hyperici) was attributed to the presence of quercetin (Schimmer *et al.*, 1988). Moreover, the mutagenic potential of the extracts was shown to correlate well with their quercetin content.

The genotoxicity of *G. macrorrhizum* extract *in vivo* was evaluated using the somatic mutation and recombination test (SMART) in *D. melanogaster*, employing flies with an increased cytochrome P450-dependent biotransformation capacity. This so-called high bioactivation cross (HB cross) makes the wing spot test more sensitive for the detection of promutagens and procarcinogens (Graf & van Schaik, 1992). The SMART data showed that at the concentrations tested the extracts did not induce somatic mutation or recombination in the *D. melanogaster* crosses. Taking into account that the SMART assay records different genotoxic events, either mutation (point mutations, specific types of chromosome aberrations, etc.) or mitotic recombination, the negative results obtained in this work indicate that *G. macrorrhizum* extracts are not mutagenic in *Drosophila*.

**Conclusions.** It seems that the protective effects of *G. macrorrhizum* extracts against <sup>1</sup>O<sub>2</sub> and their prooxidant toxicity are mainly determined by the action of quercetin and its derivatives. The data also show that the extracts from *G. macrorrhizum* are genotoxic in cytogenetic tests *in vitro*, though they revealed no genotoxic activity when investigated in the *D. melanogaster* assay *in vivo*. Our results

suggest that further investigations are necessary to determine whether the *G. macrorrhizum* extract or its components can be used without hazard to human health.

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