

Hydroalcoholic and polyphenolic extracts of *Ziziphus jujuba* mill fruits prevent methyl methanesulfonate-induced DNA damage in HepG2 cells

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Abstract

Ziziphus jujuba Mill (ZJ), which has been extensively used by the Iranian traditional healers, belongs to the *Rhamnaceae* family. This semitropical herb contains large quantities of polyphenols and flavonoids, which in turn reveal antioxidant, antibacterial, free radical scavenging, and several other pharmacological activities. The purpose of the present study was to evaluate the DNA damage prevention potential of hydroalcoholic and polyphenolic extracts of *Ziziphus jujuba* on HepG2 cells. Throughout the assessment of genoprotective properties, cells were incubated with various concentrations of hydroalcoholic (0.1, 1, 10, and 50 µg/ml) and polyphenolic extracts (0.1 and 1 µg/ml) for a one-hour period, followed by a one-hour incubation period with genotoxic concentration of methyl methanesulfonate (MMS) (10 µM). The comet assay method was applied because of its being attributable to the substantial sensitivity, its inexpensiveness, and its straightforward procedure of use. The tail length, percentage of DNA in tail, and tail moment were measured. Statistical analysis revealed that concentrations of 10 µg/ml hydroalcoholic and 1 µg/ml polyphenolic extract were genoprotective against MMS. Therefore, our results suggest that *Ziziphus jujuba* at suitable doses can prevent DNA damage.

Keywords: *Ziziphus jujuba*, genoprotective effect, HepG2, Comet Assay.

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Introduction

Constantly, cells undergo destructive oxidative stress processes, which could originate either from exogenous or endogenous sources. Cells withstand this occurrence through engaging several distinct defense mechanisms ranging from free radical scavengers such as glutathione (GSH), vitamins C, and antioxidant enzymes to the elaborate DNA repair systems (1). Genetic alterations to the DNA both in somatic and germ cells can induce a plethora of events, from single-point

mutations to chromosomal break, rearrangements or loss, which finally lead to several genetic-related diseases among which is also cancer (2). In order to identify genotoxic molecules, researchers perform specified assays contributing DNA damage in toxically affected cells. Loss of excision repair, cross-linking, alkali-labile sites, point mutations, structural and numerical chromosomal aberrations, and the compromised integrity of the genetic material have been known as the leading

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causes of carcinogenesis (3). In line with the results of preceding assessments, following a healthy diet guarantees genomic stability (4). Biomolecules found in fruits and vegetables might play a protective role in several conditions such as cancers. Revealing safer profiles has placed the naturally occurring antioxidants in a more pleasurable position than synthetic ones. Thus, the correlation between the consumption of fruits and vegetables and decreased vulnerability to cardiovascular diseases and cancer has been manifested by the existing epidemiological evidence. On account of this fact, the evaluation of genoprotective activity of fruits and vegetables has substantially attracted attentions, and a large bulk of research had been carried out (4-6).

One such medicinal plant is *Ziziphus jujuba*, which belongs to the Ramnaceae family and contains several flavonoids such as querarin, 6"-feruloylspinosin, apigenin-6-C-b-d-glucopyranoside, spinosin 6"-feruloylisopinosin, isospinosin, and isovitexin-200-O-b-d-glucopyranoside (7, 8). Additionally, a large number of phytochemicals such as 64 alkaloids, 16 glycosides, and 14 terpenoids also can be found with certain proportions in this plant. Among them, phenolic components, flavonoids, triterpenic acids, and polysaccharides are considered as efficacious radical scavengers which could exhibit potent antioxidant, anticarcinogenic, and anti-inflammatory activities which finally lead to maintaining a more stable genomic content (4, 9, 14).

Each ingredient has unique and multifactorial properties, one of the most prominent of which is its ability to limit or slow down oxidative stress reactions (9). Different parts of *Ziziphus jujuba* have been established to provide a wide range of therapeutic properties such as carminative, expectorant, and antidiabetic. In traditional

medicine, fruits and seeds of *Ziziphus jujuba* have been widely used as tonic and aphrodisiac and sometimes as sedative-hypnotic, anxiolytic, antiulcer, anti-inflammatory, antispastic, antifertility/contraception, antinephritic, cardiotonic, immunostimulant, and wound healing properties (9-14).

Assessing both genotoxicity and subsequent carcinogenicity as well as analyzing the defense and opposing protocols are fundamental prerequisites for the evaluation of pharmaceuticals. A number of fully defined *in vitro* and *in vivo* testing techniques are available in order to measure genotoxic and carcinogenic phenomenon (2).

The comet assay or SCGE (single cell gel electrophoresis) is a simple, sensitive, versatile, feasible, and economical genotoxicity measuring procedure and a specific process to detect deoxyribonucleic acid strand breaks in individual eukaryotic cells. The number of studies applying this method rises every year (15, 16). Indeed, the International Workshop on Genotoxicity Testing regards the *in vitro* SCGE in the 3-D skin model as a valuable support to genotoxicity identification as: (a) it is independent of cell proliferation, and (b) it covers a broader spectrum of DNA damage. Ostling and Johanson developed the new method of quantifying DNA damage and repair with the goal of utilization in cellular studies which imports the word

name implied data was analyzed by a software (Comet score) to provide the protocol with a higher definition sensitivity (2, 17, 18).

Methyl methanesulfonate (MMS) is the most common methylating agent which, for years, has been applied as an experimental research chemical. MMS acts through modification in both guanine and adenine to create base mispairing and replication

blocks, respectively. DNA-related destructions caused by alkylating agents would principally be retrieved via the base excision repair (BER) pathway and DNA alkyltransferase actions (19, 20). In the present study, we used HepG2 cells (hepatoma cells) for specialized liver function and comparable activities with human hepatocytes (21).

Based on the aforementioned research findings, this research aimed to investigate the genoprotective activities anticipated from hydroalcoholic and polyphenolic extracts of *Ziziphus jujuba* on HepG2 cells against MMS toxicity by taking advantage of the SCGE method. Additionally, according to each extract, the genotoxic features would be specified individually. And, the correlation between DNA damage quantity and various concentrations of this herbal extract would be mentioned illustratively.

Materials and Methods

Materials

Ethanol, Chloroform, EDTA, H₂O₂, NaCl, NaOH, Na₂CO₃, NaH₂PO₄, Folin Ciocalteu reagent (FCR), Tris, and Triton X-100 were acquired from Merck Co. (Germany). Low melting point agarose (LMA), Na₂HPO₄, KCl, methyl methanesulfonate (MMS), and ethidium bromide were from Sigma Co. (USA). Normal melting point agarose (NMA) was supplied by Cinnagen Co. (Iran). The RPMI 1640 medium and fetal bovine serum (FBS) were obtained from Capricorn (Austria), and antibiotic was purchased from PAA Co. (Austria). And, HepG2 cells came from Pasture Institute (Iran). The *Ziziphus jujuba* fruits were purchased from a local medicine market in Isfahan (Isfahan province, Iran) in November 2013 and authenticated by Pharmacognosy Department of Isfahan Pharmacy School as *Ziziphus jujuba*. A sample of plant is deposited in our school

for future evidence (Herbarium number: 1893).

Extract Preparation

The *Ziziphus jujuba* fruits were carefully cut into slices. The small cut slices were air-dried at room temperature, then powdered, and extracted with 500 ml of 70% aqueous EtOH through maceration (48 hours for two times) at room temperature to obtain hydroalcoholic extract. Then, all the extract was evaporated in a rotating evaporator and was freeze-dried. Pulverized pulp (100 g) of *Ziziphus jujuba* was extracted in two steps; first, with EtOH:H₂O with a proportion of 90% (v/v) and then with EtOH:H₂O with a proportion of 50% (v/v). At each step, a sufficient amount of solvent was added to make liquid slurry, and the mixture was left for 12 hours. The resultant product was filtered by Büchner funnel under vacuum condition and then was freeze-dried. The two extracts were then combined and evaporated to 1/3 of the original volume. The resultant aqueous extract was first cleared by extraction (in a separating funnel) with chloroform and was then evaporated to be completely dried under vacuum in a rotary evaporator and freeze-dried (22-24).

Cell Culture

In the present research, human hepatoma (HepG2) cells were applied to find the genotoxicity and genoprotective effect of *Ziziphus jujuba*. The HepG2 cell line was obtained from Pasture Institute of Iran and was maintained in RPMI medium (containing 7% fetal bovine serum and 1% penicillin/streptomycin to avoid the growth of unwanted and pathogenic bacterial microorganisms) and was incubated in a humidified atmosphere of 95% air in micro filter plates. To select the lowest genotoxic dose, HepG2 cells were incubated with

different concentrations of MMS, for 1 hour. A series of concentrations in the

diluting the stock solution (11.82 M) with RPMI. Then, Cells were incubated with different concentrations of extracts followed by a one-hour period of incubation with selected concentration of MMS in 12-wells plates. After that, the upper medium of wells was thrown away and washed with PBS. After trypsinization, 1 ml of medium was added to each falcon tube to be utilized for the next stages of the SCGE. For implementing SCGE, at least 90% of viability was required (24-26).

Alkaline Comet Assay

The comet assay (SCGE) is a gel electrophoresis method that is used to visualize and measure DNA strand breaks in individual cells using microscope. In its simplest form, incubated cell suspensions (0.25×10^6 cells per well) were mixed with 1% LMP agarose (37 °C). Then, they were placed on the precoated slides (1% NMP agarose). Afterwards, they were embedded on a microscope slide, and they were covered by cover glasses for 5 minutes and then they were immersed in a lysis solution (pH = 10.0) for 40 minutes and rinsed with distilled water to remove lipids and proteins. To run the electrophoresis protocol, the slides were put in a weak buffer (pH > 13.0, for 40 minutes) in order to separate broken DNA. After electrophoresis was done, 25 V with an electricity current adjusted to 300 mA, DNA was stained using a fluorescent dye 5 minutes, then washed in ice-cold medium or phosphate buffered saline, and viewed using under $\times 400$ magnification using a fluorescence microscope with an excitation filter of 510-560 nm and barrier filter of 590 nm. All stages of SCGE were performed at dark conditions, and all

solutions were prepared freshly and used cool. Individual images can then be digitized and analyzed for informative properties such as the distance that the DNA has migrated and the percentage of DNA that has migrated. These features give an indication of the number of strand breaks present in the cell (16, 24, 25, 27, 28).

Folin-Ciocalteu method for measurement of total phenolic content

20 μ L extract 5g/l was added to 1.58 mL deionized water into test tubes and 100 μ L Folin-Ciocalteu reagents were mixed. Then 300 μ L of 20% sodium carbonate solution was added to the mixtures. The tubes were mixed, and then were allowed to incubate for 2 h at 20 °C. The absorbance of the resulting blue color was measured by colorimetric at 765 nm. A calibration curve of Gallic acid (ranging from 50 mg/l to 500mg/l) was prepared (31).

Statistical Analysis

Based on this research, the most common parameters analyzed are tail length (the length of the comet tail), the percentage of DNA in the tail (percentage of colored spots in tail), and tail moment (percentage of DNA in tail \times tail length). The percentage of DNA in the tail is generally defined as the amount of DNA in the tail divided by the amount of DNA in the cell multiplied by 100, while the tail length is the distance from the middle or the estimated perimeter of the comet head to the last visible signal in the tail. These factors were used for statistical analysis in this investigation (15, 29). The raw data that were obtained by comet score software were imparted as means \pm SD (standard deviation), three continuous quantitative variables (comet length, percentage of DNA in tail, and tail moment) were analyzed, and post hoc test (Tukey) was performed on raw data to determine

significant differences between concentrations. Statistical analysis was accomplished by IBM-SPSS software, USA.

Results

In order to determine protective effects of hydroalcoholic and polyphenolic extracts of *Ziziphus jujuba* on genotoxicity induced by MMS, HepG2 cells were incubated with different concentrations of extracts for 1 hour followed by a one-hour incubation period for MMS (10 μ M). The results of SCGE after the one-hour incubation period for both extracts were analyzed. One-way analysis of variance (ANOVA) for the tail length extracts was significant ($p < 0.001$). According to the results of Tukey's multiple comparison post hoc test, all concentrations have decreased the tail length significantly ($p < 0.001$) in comparison with the MMS group. The results of one-way ANOVA for the percentage of DNA in tail were significant ($p < 0.001$). Based on the results of the Tukey's multiple comparison post hoc test, in all concentrations of both extracts the percentage of DNA in tail decreased significantly ($p < 0.001$) compared to the MMS group. The result of one-way ANOVA of the tail moment for all groups was significant ($p < 0.001$). Moreover, the results of Tukey's multiple comparison post hoc test for all concentrations showed a significant decrease ($p < 0.001$) in this parameter compared to the MMS group.

SCGE Results of Different Concentrations of MMS

In order to determine genotoxic concentration of MMS, HepG2 cells were incubated with different concentrations of 1, 5, 10, 100, and 500 μ M for 1 hour followed by the SCGE. Regarding the results of this step, the best concentration of MMS (10 μ M) had significant difference with the control group (Fig. 1). In addition,

the number of incubated cells showed no decrease during incubation. Hence, this concentration was selected for the next stage of the study. At first, the minimum amount of genotoxic concentration of MMS was determined. The statistical outcomes showed significant differences between concentrations of 10, 100, and 500

($p = 0.05$) was selected as the minimum amount of genotoxic concentration of MMS.

Studying Genotoxic and Genoprotective Effects of Extract of ZJ Combined with MMS

Three repetitions of SCGE were performed for each extract concentration and analysis was implemented on at least 100 cells per slide. To determine the genotoxic effects, concentrations were contrasted with negative control, and significant differences were manifested by (*) sign.

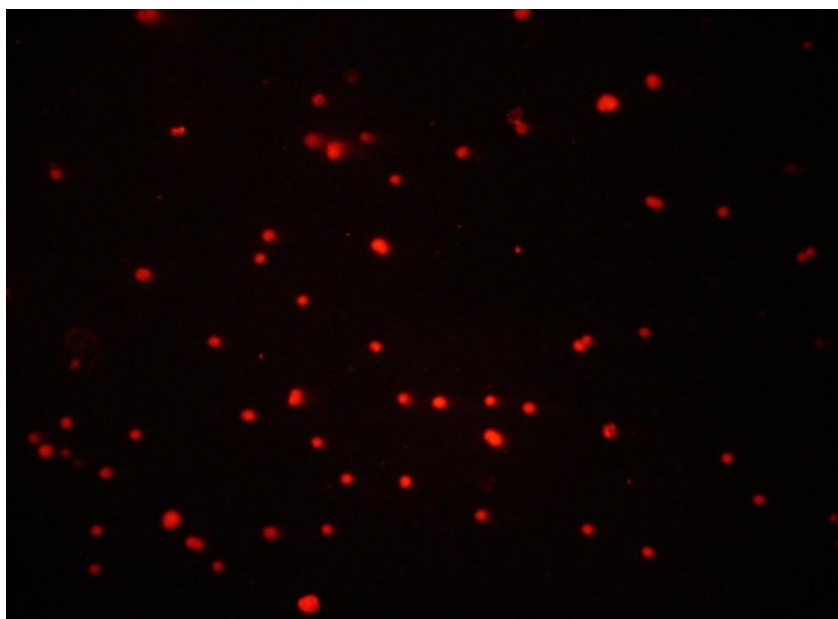
Subsequently, genotoxic effects of hydroalcoholic extract (data not shown) were surveyed in various concentrations (1, 10, 100,

concentrations and negative control, and drastically different from negative control

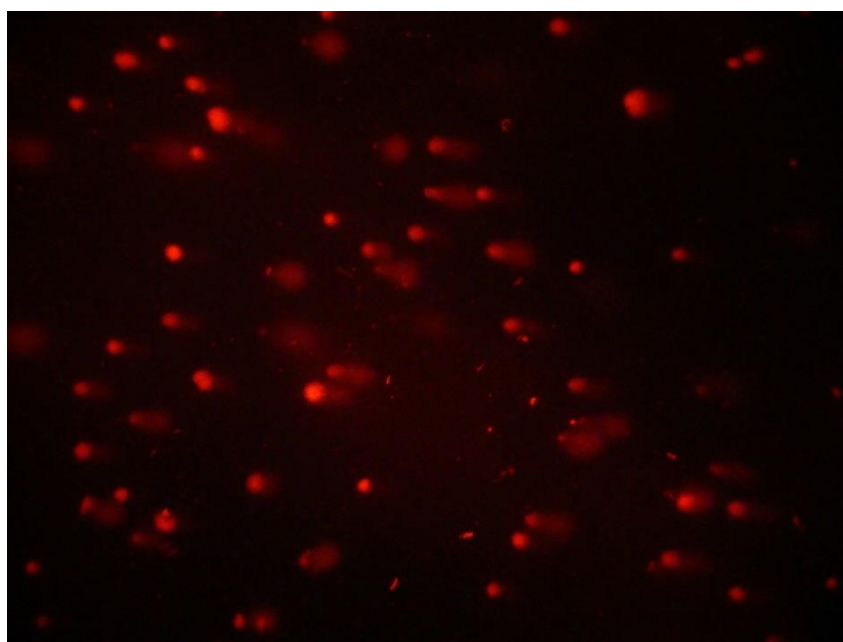
and lower concentrations were safe and had no genotoxic effect (Table 1). Consequently, genoprotective effects of hydroalcoholic extract were determined in 1, 10, and 50

cellular exposure was performed with pre-incubation of cells with these concentrations for 1 hour. Then, HepG2 cells were incubated by minimum genotoxic concentration of MMS as

Results manifested significant differences



A



B

Figure 1 A) Comet image of untreated HepG2 cells as negative control. B) Comet image of HepG2 cells treated with 10 μ M of MMS as positive control

Table1 Genotoxic effects of *Ziziphus jujuba* Mill extracts on HepG2 cells

Treatment	Concentration (µg/ml)	Tail length(pixels) (Mean ± SD)	%DNA in tail (Mean ± SD)	Tail moment(pixels) (Mean ± SD)
<i>Hydroalcoholic extract</i>	Negative control (PBS)	0.56 ± 0.03	0.09 ± 0.14	0.0007 ± 0.002
	1	0.96 ± 0.37	0.27 ± 0.81	0,007 ± 0.035
	10	4.4 ± 1.1	2.5 ± 0.8	0.3 ± 0.27
	100	60.9 ± 0.6*	22.8 ± 2.1*	18.8 ± .6*
	500	98 ± 2.7*	22 ± 1.7*	29 ± 2.6*
<i>Polyphenolic extract</i>	Negative control (PBS)	1.64 ± 0.18	0.46 ± .94	0.02 ± 0.09
	1	1.52 ± 0.6	0.97 ± 0.18	0.03 ± 0.04
	10	51.7 ± 2.6*	24.9 ± 2.9*	14.4 ± 2.2*
	100	108 ± 6.9*	38 ± 2*	44 ± 4*
	500	131 ± 6.2*	52 ± 1.2*	70 ± 1.3*

The effect of genotoxic concentrations of hydroalcoholic and polyphenolic extract of *Ziziphus jujuba* (µg/ml) in comparison with negative control on tail length (pixels), %DNA in tail, and tail moment (pixels) of 3 independent experiments are represented as mean ± SD. Significant difference ($p < 0.001$) was illustrated by (*) sign.

µM of MMS, and mentioned concentration had genoprotective effects on MMS (0.001).

After this step, genotoxic effects of various concentrations of polyphenolic extract (1,

Therefore, a meaningful difference was

negative control, and the lower concentrations were safe and had no genotoxic effect (Table 1). Afterwards,

genoprotective effects of this extract were determined by various concentrations (0.1, 1 concentrations had genoprotective effects consistent with positive control and the 1 protection (Table 2).

Total phenolic content of extracts

According to the Folin-Ciocalteu method, total amount of phenolic content assayed, and total phenolic content of the

Table 2 Genoprotective effects of *Ziziphus jujuba* Mill on HepG2 cells using the comet assay

Treatment	Concentration range ($\mu\text{g/ml}$)	Tail length (pixels) (Mean \pm SD)	%DNA in tail (Mean \pm SD)	Tail moment (pixels) (Mean \pm SD)
<i>Hydroalcoholic extract</i>	Positive control (MMS)	107 \pm 3.8	39 \pm 2.6	46 \pm 6.1
	1	8.9 \pm 1.1*	11 \pm 1.3*	2.4 \pm 0.04*
	10	6.8 \pm 0.8*	6.5 \pm 0.5*	0.8 \pm 0.3*
	50	85 \pm 0.5	28.7 \pm 0.8	28.8 \pm 2.9
<i>Polyphenolic extract</i>	Positive control (MMS)	107 \pm 5.3	49 \pm 2.5	57 \pm 5.1
	0.1	104 \pm 6.1	33 \pm 2.7	39 \pm 5.3
	1	8.4 \pm 0.1*	7.5 \pm 6.8*	1 \pm 0.4*

The effect of genoprotective concentrations of hydroalcoholic and polyphenolic extract of *Ziziphus jujuba* compared with control groups on tail length (pixels), percentage of DNA in tail, and tail moment (pixels) that are represented as mean \pm SD. * Mean value was significantly different from control (01) one-way ANOVA followed by post hoc test).

hydroalcoholic and polyphenolic extracts was reported to be 1.8% and 2.8% of Gallic acid, respectively. All experiments were performed in triplicate.

Discussion

The objective of this cellular toxicity analysis was to assess the distinct concentration ranges of *Ziziphus jujuba* extracts in order to demonstrate whether they possess either genoprotectivity or

genotoxicity features. Different parts of *Ziziphus jujuba* fruit could be considered as a rich source of antioxidant and antiproliferative compounds including cyclopeptide alkaloids, flavonoids, sterols, jujuboside A, jujuboside B, lauric acid, triterpenoid saponins, ascorbic acid, anthocyanines, and polyphenolic components which play roles in genomic stability and in the reduction of internal or external oxidant. Therefore, it is supposed

to be an appropriate preventive agent against cancerous disorders besides any free radical-related body dysfunction (9, 10, 30, 31). As illustrated by the result of the present study, it is noteworthy to mention that *Ziziphus jujuba* extract would be genoprotective and genotoxic for the cell DNA, which depends on the concentration and the type of solvent used. Some extracts cause DNA damage and induce cell death and show anti-cancer effects (32). In accordance with the results, and also attending the SCGE method accuracy and sensitivity as well as selected concentration (each of these could be an error source and regarded as limitations for the present study), hydroalcoholic and polyphenolic extract had shown genotoxic effect in 100

concentrations below the genotoxic ones, there are biologically active components in hydroalcoholic extract that exhibited dose-related protective behavior against MMS-induced DNA damage where 1 and 10 reduction in tail length, the percentage of DNA in tail, and tail moment comparing to positive control. The

probably MMS genotoxicity stands over the protective effect of this specific herb. This could be deduced in two rationalities as antioxidant capacity reduction or genotoxic properties commence. Based on our data,

regarded genotoxic. Therefore, genomic protective ability was analyzed in 1 and 0.1 table 2

indicates, the best concentration for producing genoprotective aspects is

significantly different from positive control.

Conclusion

When comparing these two extracts, due to the higher antioxidant and genoprotective ability revealed in lower concentrations by the polyphenolic solution, it would be regarded as a more potent complex against genomic content instability.

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Conflict of interest

The authors declare that they have no conflict of interest.

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