

THE ANTIMUTAGENIC EFFECT OF MULTIFLORAL HONEY IN SALMONELLA/ MICROSOMAL ASSAY AND ITS CORRELATION WITH THE TOTAL POLYPHENOLIC CONTENT

Mabrouka BOUACHA^{1,*}, Ines BOUDIAR², Akila ABDI², Mohammad Abdulraheem Al-KAWAWEEN³ and Messaouda KHALLEF⁴

Address(es): Mabrouka BOUACHA

¹Laboratory of Biochemistry and Environmental Toxicology, Department of Biochemistry, Faculty of Sciences, University of Badji Mokhtar, Annaba, Algeria.

²Laboratory of Biochemistry and Microbiology, Department of Biochemistry, Faculty of Sciences, University of Badji Mokhtar, Annaba, Algeria.

³Faculty of Pharmacy, Department of Pharmacy, Al-Zaytoonah University of Jordan, Amman, Jordan

⁴Department of Biology, Faculty of Natural Sciences, Life Sciences, Earth and the Universe, university of 8 Mai 1945, Guelma, Algeria

*Corresponding author: mabrouka.bouacha@univ-annaba.dz

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ABSTRACT

The objective of this study is to determine the antimutagenic effect of honey and its correlation with the concentration of the total polyphenolic content. Seven honey samples were collected from different regions of Algeria. The total polyphenolic content was determined by Folin-Ciocalteu colorimetric assay. The antimutagenic effect was carried out by the AMES *Salmonella*/microsome mutagenicity assay against three known mutagenic substances (4-nitro-*o*-phenylenediamine, sodium azide, and mitomycin C), using *Salmonella typhimurium* TA98, TA100, and TA102 strains.

The results obtained revealed that Algerian honey contains high polyphenol content, which varied significantly between 38.04 and 286.28 µg of GAE/100 mg of honey. This variation is due to their different botanical and regional origins. In addition, all tested honey exhibited an antimutagenic effect against mutagenic substances; honey is effective to inhibit between 29.18±11.11 % and 73.14±11.14 % of mutagenic activity of chemical substances. There is a strong positive correlation between the total polyphenolic content and the antimutagenic properties of honey against the mutagenic substances.

The results obtained suggest that honey is effective as an antimutagenic agent; it can play an important role in the protection of the mutagenic effect of DNA caused by chemical substances.

Keywords: AMES assay, antimutagenic effect, honey, phenolic content, *Salmonella typhimurium*

INTRODUCTION

Honey is a natural product containing several nutritional and therapeutic properties that are gained from the nectar of flowers (Nikhat and Fazil, 2021; Al-kafaween et al., 2021). It has been used since ancient times as a flavorful sweetener and it is considered a part of traditional medicine (Wang, Andrae and Engeseth, 2002; Alotibi et al., 2018). It consists of a highly concentrated solution of a complex mixture of sugar and other constituents, such as minerals, proteins, vitamins, organic acids, flavonoids, phenolic acids, enzymes, and volatile compounds (Alvarez-Suarez et al., 2009; Sachdev, Kumar and Ansari, 2021). Honey presents a wide range of biological effects such as antimicrobial (Bouacha, et al., 2018; Bouacha and Benbouzid, 2020), wound healing (Alvarez-Suarez et al., 2014; Febriyenti et al., 2019), antioxidant (Ahmed et al., 2018; Erejuwa et al., 2012), anti-inflammatory (Liao et al., 2018), gastroprotective (Djebli et al., 2021), and anticancer properties (Abbas et al., 2021). However, the quality of honey depends strongly on the source of nectar, plant compounds, geographical region, seasonal and environmental factors, and storage conditions (Ahmed et al., 2018; Matzen et al., 2018).

The compounds that are responsible for the major biological effects of honey are the total polyphenolic content (Gómez-Maqueo et al., 2018; Rodrigues da Silva, Campos Chisté and Fernandes, 2021). Recently, the total polyphenolic content in diet has received increasing attention due to some interesting discoveries about their therapeutic properties (Becerril-Sánchez et al., 2021; Celeiro et al., 2021; González-Ceballos et al., 2021). Indeed, the total polyphenol content such as phenolic acids, naphthoquinones, xanthenes, stilbenes, flavonoids, lignans, lignins, and condensed tannins have become more and more important for scientists who are searching for novel compounds capable to prevent several human diseases, including cancer diseases. This is due to their antimutagenic and anticarcinogenic properties (Belščak-Cvitanović et al., 2018; Gómez-Maqueo et al., 2018; Santos-Buelga et al., 2019).

Many mutagenic agents are known as direct-acting mutagens, such as sodium azide (NaN₃), which directly affect genetic material that can cause point mutation in the genome and structural damage. The 4-nitro-*o*-phenylenediamine can indirectly act on DNA via the induction of the synthesis of different chemicals which can affect DNA directly and causes frameshift mutations (Słoczyńska et al., 2014). Mitomycin C, an antibiotic used in therapy against many forms of human cancer, can interact with biological molecules and can induce genetic hazards in non-tumor cells (Hammarsten et al., 2021). One of the possible approaches to protect DNA from damage caused by mutagen agents is to use natural antimutagenic products that can counteract the effects of mutagenic substances. Hence, searching and identification of antimutagenic and anticarcinogenic components represent a rapidly expanding field of cancer research (Słoczyńska et al., 2014).

Algeria, found in the northern region of the African continent, exhibits highly diversified forest ecosystems and a significant variation in climate that varies from the Mediterranean to the Saharan type. These conditions offer diverse types of high-quality honey that are rich in phytochemical bioactive compounds. Up to date, very few studies have been conducted to investigate the properties and application of Algerian honey in human diseases. Indeed, to the best of our knowledge, this is the first study that reports the antimutagenic effect of honey and its relationship with the total polyphenolic content. Therefore, we sought to determine the antimutagenic effect and the possible preventive capacity of honey from seven different floral and geographic sources of Algeria against the mutagenicity of three positive mutagens: 4-nitro-*o*-phenylenediamine, sodium azide, and mitomycin C, and its relationship with the total polyphenolic content of honey.

MATERIAL AND METHODS

Honey samples

Seven Algerian multifloral honey samples from *Apis mellifera* were obtained directly from beekeepers from different geographical regions and vegetation

Table 1 Botanical and geographical locations of the harvested honey

Honey samples	Floral source	City	Geographical region
1	<i>Eucalyptus, Pinus</i>	El Taref	Extreme north-eastern of Algeria
2	<i>Quercus, Castanea</i>	Jijel	North-eastern of Algeria
3	<i>Citrus:C.maxima, C. sinensis, C. aurantifolia, C. limon</i>	El Blida	The central part of the north of Algeria
4	<i>Thymus hirtus, Marrubium vulgare</i>	M'Sila	The central part of the north of Algeria
5	<i>Ziziphus, Artemisia,</i>	Djelfa	Central part of north of Algeria
6	<i>Ruta graveolens, Pituranthos scoparius,</i>	Batna	East of Algeria
7	<i>Rosmarinus officinalis, Ecballium elaterium, Lonicera caprifolium</i>	Tebessa	East of Algeria

Tester strains

The *Salmonella typhimurium* tester strains used in AMES assay were unable to grow in the absence of histidine because each tester strain contains a different mutation in various genes in the histidine operon; this makes the tester strains unable to synthesize this amino acid. The tester strain *Salmonella typhimurium* TA98 was used to detect mutagenic substances causing frameshift mutations. However, TA100 was used to detect mutagens causing base-pair substitutions mutations and TA102 was used to detect mutagens causing transition/transversion mutations (Mortelmans and Zeiger, 2000; Tarawneh et al., 2021; Al-Bakri et al., 2019; Huwaitat et al., 2021).

Determination of total polyphenolic content

The determination of total phenolic content was performed using the Folin-Ciocalteu method (Deng et al., 2018), and the results were expressed as µg gallic acid/mg of honey. Initially, 0.5mL of honey solution was mixed with 0.3 mL of the Folin-Ciocalteu reagent and 2 mL of a 15% sodium carbonate solution. Distilled water was added to a final volume of 5 mL. All samples were incubated at room temperature in the dark conditions for 2 hours, and their optical density was read at 760 nm against a blank of distilled water. A standard curve of gallic acid was drawn within a concentration range of 7.0×10^{-4} to 7.8×10^{-3} mg/mL. The linearity obtained was 0.982 (Chaiyasut et al., 2018).

Antimutagenic effect

The antimutagenic effect of honey was achieved according to the lightly modified assay described previously by Mortelmans et Zeiger, (2000). This assay evaluates the ability of a chemical or a physical agent to cause specific mutations in different strains of *Salmonella typhimurium*. These strains are carrying a specific mutation in one of the genes encoding the synthesis of histidine. This mutation makes the bacteria unable to grow on a medium without histidine (His⁻). By exposing the His⁻ bacteria to mutagenic agents; the histidine (His⁻) mutations reverse to His⁺.

In this study, three strains of *Salmonella typhimurium* with three positive mutagens were used: 4-nitro-o-phenylenediamine for *S. typhimurium* 98, sodium azide for *S. typhimurium* 100, and mitomycin C for *S. typhimurium* 102. The presence of genetic markers and the plasmids (pKM101 and pAQ1) has been systematically checked.

Each bacterium was cultured in 20 mL of nutrient broth and incubated for 24 h at 37 °C with continuous agitation. 100 µL of overnight culture, 100 µL of each honey, and 100 µL of positive mutagen are mixed with 500 µL of phosphate buffer (13.8 g/L NaH₂PO₄ and 14.2g/L Na₂HPO₄). The mixture is pre-incubated at 37°C for 20 minutes.

A volume of 2 mL of molten soft agar medium with histidine and biotin solution was added aseptically to the mixture and poured after slight agitation into a minimal glucose agar plate. After solidification of the mixture, the plates are incubated for 48 hours, at 37°C. The number of revertant colonies on the plates was counted after incubation (Mortelmans and Zeiger, 2000; Al-kafawenn et al., 2021).

Honey is considered to have a mutagenic effect if the reversion coefficient (RC) is greater than or equal to two. This coefficient is calculated according to this formula: $RC = \frac{R_1}{R_0}$

The antimutagenic effect of honey is expressed as a percentage of inhibition (I %) of mutagenicity of chemical substances, it is calculated according to the following formula: $I (\%) = 100 - \left(\frac{R_3}{R_2} \times 100 \right)$

Where: R₀ is the number of spontaneous revertant colonies per plate
 R₁ is the number of revertant colonies induced by honey
 R₂ is the number of revertant colonies per plate exposed to positive mutagen
 R₃ is the number of revertant colonies per plate exposed to positive mutagen and honey.

sources (Table 1). Honey samples were collected in a sterile container and kept at 4 °C in darkness until usage. The botanical origin of the honey samples was directly related to the type of plants that the honeybee gleaned. Honey samples were collected in a sterile Erlenmeyer flask and the samples were stored at room temperature and were hidden from the light exposure until usage.

Statistical analysis

Significant differences were calculated using analysis of variance (ANOVA) followed by a multiple ranges test (Tukey). Correlations between the antimutagenic effect of honey and total polyphenolic content were obtained using correlation coefficient (r). All analyses were carried out using the software GraphPad Prism version 7.00 (Graph Pad Software, Inc., La Jola, CA, USA).

RESULTS AND DISCUSSION

Determination of total polyphenolic content

The total phenolic content of Algerian honey represented (Figure 1) was found to vary significantly, between 38.04 and 286.28 µg of GAE/100g of honey. This range is similar to the ranges typically found for European (Kuś et al., 2014) and African honey samples (Beretta et al., 2005). While, these values are also considered relatively high in comparison with those found in other regions in Algeria (14.50 to 99.62 µg of GAE / 100 mg) or other countries such as Brazil (26.00 to 100.00 µg of GAE/100 mg of honey) (Nascimento et al., 2018; Zerrouk et al., 2018). Ciappini and Stoppani (2014) reported that the total polyphenolic content in Argentinean honey varied from 40.30 to 193.00 µg of GAE /100 mg of honey. Indeed, the floral source and geographical region of honey samples mainly influence the type and concentration of polyphenolic content; these findings are in agreement with results reported previously (Al et al., 2009; Ciappini and Stoppani, 2014; Cianciosi et al., 2018; Cheung et al., 2019).

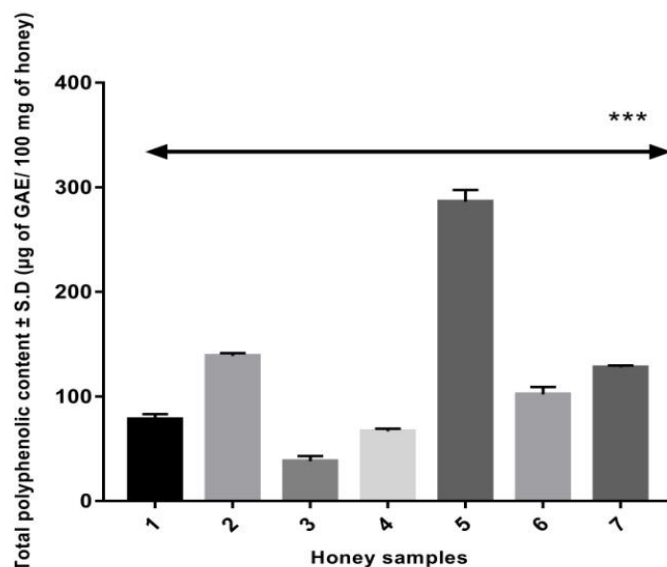


Figure 1 Concentration of total polyphenolic content in honey (µg of GAE/100 mg of honey), calculated as the mean of three repeated experiments ±SD. ***: means that there are very highly significant differences between the total polyphenolic content of honey samples at P<0.0001

Mutagenic and antimutagenic effect of honey

As shown in Table 2, none of the honey samples showed a mutagenic effect, hence honey is considered as a non-mutagenic product. However, all honey samples have displayed a significant antimutagenic effect against the three positive mutagens.

The percentage of antimutagenicity is ranged from 29.18±11.11 to 73.14±11.14%. Honey is more effective to reduce the number of revertant colonies in *S.typhimurium* TA98 and TA100 than in strain TA102, and this can be explained by the type of mutations. Since strains TA98 and TA100 are known to detect frameshift and base-pair substitution mutation types, respectively, strain TA102 is known to detect transition/transversion mutation types. In addition, the results showed that honey samples 5 and sample 2 can reduce the highest percentage of revertant colonies on the three tested strains. These results are in concordance with those of Saxena et al. (2012); reported that honey exhibits highly inhibition of mutations caused by ethyl methanesulfonate, ultraviolet, and gamma radiation in *S.typhimurium* TA100 and TA102. Honey is effective to reduce the mutagenicity induced by 4-Nitro-*o*-phenylenediamine, sodium azide, and mitomycin. There are highly significant differences among the antimutagenic effects of honey

($P<0.0001$). It appears that honey has more antimutagenic effect in *S. typhimurium* TA98 and TA100 strains than in *S. typhimurium* TA102 strain; this is probably related to the tester strains and to the differences in mechanistic pathways of the antimutagenic effects of honey. The percentage of inhibition of mutagenicity varied from 26.18 to 73.14%. This range is comparable to the range found by Saxena et al. (2012) which is between 3 and 77%. Indeed, honey was found to be highly effective to inhibit frameshift and base-pair substitution mutation types than transition/transversion mutation types. Very few studies were reported regarding the potential antimutagenic effects of honey in bacterial strains; however, Wang et al. (2002) have reported that floral honey from the USA exhibits antimutagenicity against Trp-p-1, a heterocyclic food mutagen in Ames Assay.

Table 2 Mutagenic and antimutagenic effect of seven honey on *Salmonella*/microsomal assay represented as a revertant coefficient (RC±SD) and percentage of inhibition of mutagenicity (I±SD %), respectively.

Honey samples	<i>S. typhimurium</i> TA98		<i>S. typhimurium</i> TA100		<i>S. typhimurium</i> TA102	
	RC±SD	I±SD %	RC±SD	I±SD %	RC±SD	I±SD %
1	1.10±0.11	50.45±07.14 ^a	1.21±0.28	49.65±07.91 ^a	1.23±0.02	29.18±11.11 ^a
2	1.09±0.25	62.26±14.11 ^b	1.23±0.09	52.81±11.32 ^a	1.17±0.07	36.43±06.34 ^b
3	1.14±0.31	49.24±09.62 ^a	1.08±0.49	44.14±12.12 ^b	1.18±0.18	26.18±09.78 ^a
4	1.21±0.27	53.17±12.84 ^a	1.27±0.45	50.81±06.85 ^a	1.22±0.27	27.12±12.04 ^a
5	1.07±0.39	73.14±11.14 ^c	1.15±0.12	68.61±14.79 ^c	1.13±0.19	38.44±08.12 ^c
6	1.32±0.28	59.43±7.28 ^b	1.19±0.32	53.01±09.43	1.14±0.32	26.52±11.67 ^a
7	1.07±0.13	57.54±05.12 ^b	1.12±0.28	57.25±04.78 ^d	1.25±0.10	32.32±12.11 ^b

Correlation between antimutagenic effect and total polyphenolic content

There is a strong positive correlation between total polyphenolic content and the efficiency of honey to inhibit the mutagenicity of 4-nitro-*o*-phenylenediamine, sodium azide, and mitomycin C (Figure 2). The correlation coefficient is ranged between 0.806 and 0.938, it was found in decreasing order as follows TA100>TA98>TA102. It is known that honey contains several bioactive substances others that total polyphenolic content such as sugar, vitamins, trace elements, amino acids, proteins and carotenes, organic acids as well as certain enzymes including glucose oxidase, invertase, and catalase (Erejuwa, Sulaiman and Ab Wahab, 2012); which may contribute in the antimutagenic properties of honey. Indeed, honey has a protective effect against mutagenic substances. These findings are in agreement with those of Meskini et al., (2018); they have proved that the topical application of honey inhibits the carcinogenic effects of 7,12-Dimethylbenz(a)anthracene- on mice skin. In addition, the results reported by Fauzi et al., (2011); showed that Tualang Honey has a significant anticancer activity, which induces apoptosis and disrupts the mitochondrial membrane potential of human breast and cervical cancer cell lines.

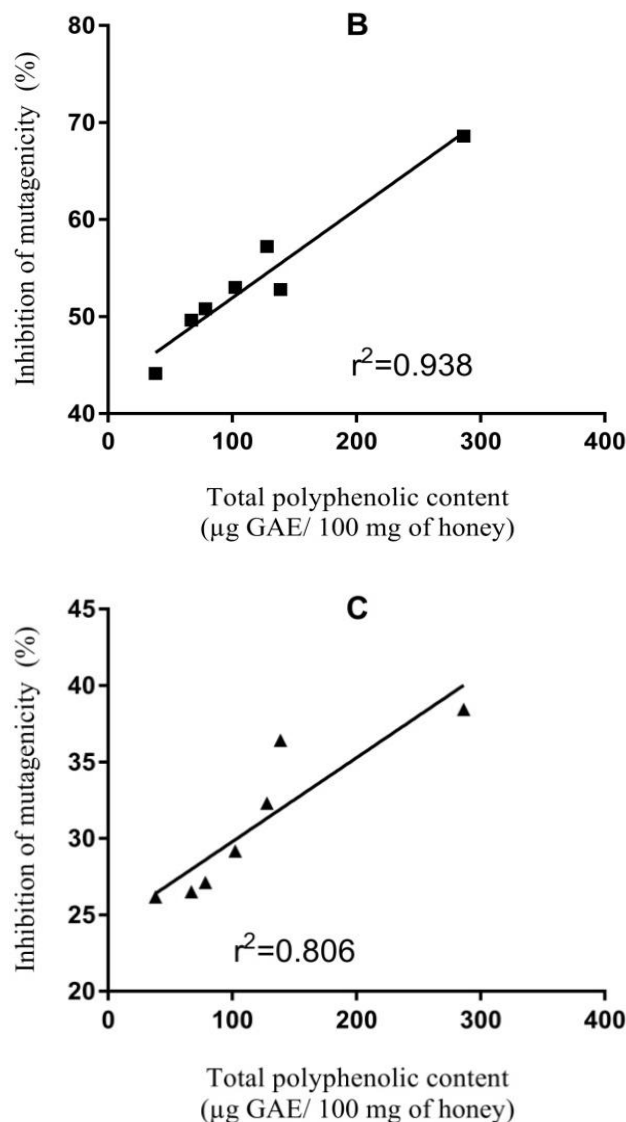


Figure 2 Correlation between the concentration of total polyphenolic content in honey and the inhibition of the mutagenic effect of chemical substances in *Salmonella typhimurium* TA98 (A), *Salmonella typhimurium* TA100 (B), and *Salmonella typhimurium* TA102 (C). r^2 : is the r square of the coefficient of correlation

The antimutagenic effect of total polyphenolic content of natural compounds such as plant extracts and vegetable food extracts was previously described by others authors (de Mejía, Castaño-Tostado and Loarca-Piña, 1999; Sloczyńska et al., 2014; Delgado-Vargas et al., 2018; Temviriyankul et al., 2021). However, the mechanisms and the types of active compounds involved in the protective effects of plants against DNA mutations have not been clearly identified. Whereas, reduction of oxidative stress is the main factor in the prevention of DNA mutations and in the decrease of chronic disease risk, including some forms of cancer (Makhafola et al., 2016; Tarawneh et al., 2021; Al-Bakri et al., 2019; Huwaitat et al., 2021). Moreover, the antimutagenic effect of honey may be attributed to different mechanisms, including cell cycle arrest, activation of mitochondrial pathway, induction of mitochondrial outer membrane permeabilization, induction of apoptosis, modulation of oxidative stress, amelioration of inflammation, modulation of insulin signaling, and inhibition of angiogenesis in cancer cells (Erejuwa et al., 2014). These findings are in agreement with those of Cianciosi et al. (2018) and Miguel et al. (2017); they declared that the phenolic compounds are associated with health benefits of honey; ranging from antioxidant, immunomodulatory, and anti-inflammatory activity to anticancer effects (Abbas et al., 2021). On another hand, we suggest in this study that the concentration of total polyphenolic content is an important indicator in the choice of honey of good quality since it correlates positively with the antimutagenic effect of honey.

CONCLUSION

In conclusion, we reported in this study that honey may act to decrease the incidence of DNA mutations; this is correlated with the phytochemical compounds in honey, particularly, the total polyphenolic content. Indeed, there is a strong correlation between the antimutagenic effect of honey and its composition with total polyphenolic content.

In addition, we suggest in this study that honey may contribute as an alternative to pharmaceutical medications that can prevent and inhibit the DNA mutations responsible for serious human diseases, including cancer. However, further research on the pathway and mechanism of action of honey in the reduction of DNA mutations is highly recommended.

Conflict of interest: We declare that we have no conflict of interest.

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