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Biocompatibility, Cytotoxicity, Antimicrobial and Epigenetic Effects of Novel Chitosan-Based Quercetin Nanohydrogel in Human Cancer Cells

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Background: Previous studies have reported that quercetin (Q) has a potential antibacterial and anticancer activity. However, its application is limited by many important factors including high hydrophobicity and low absorption.

Methodology: In the current study, we synthesized and characterized (Patent) a novel chitosan-based quercetin nanohydrogel (ChiNH/Q). Encapsulation efficiency was confirmed by UV/VIS spectrophotometer. Physicochemical characterization of ChiNH/Q was assessed by PDI, DLS, SEM, FTIR, and XRD. The toxicity of the ChiNH/Q against five strains of the pathogen and HepG2 cells was examined. Moreover, the quantification of ChiNH/Q on genomic global DNA methylation and expression of DNMTs (DNMT1/3A/3B) in HepG2 cancer cells were evaluated by ELISA and real-time PCR, respectively.

Results: Under the SEM-based images, the hydrodynamic size of the ChiNH/Q was 743.6 nm. The changes in the PDI were 0.507, and zeta potential was obtained as 12.1 mV for ChiNH/Q. The FTIR peak of ChiNH/Q showed the peak at 627 cm⁻¹ corresponded to tensile vibrational of NH_2 -groups related to Q, and it is the indication of Q loading in the formulation. Moreover, XRD data have detected the encapsulation of ChiNH/Q. The ChiNH/Q showed a potent antimicrobial inhibitory effect and exerted cytotoxic effects against HepG2 cancer cells with IC_{50} values of 100 μ g/mL. Moreover, our data have shown that ChiNH/Q effectively reduced (65%) the average expression level of all the three DNMTs (p<0.05) and significantly increased (1.01%) the 5-methylated cytosine (5-mC) levels in HepG2 cells.

Conclusion: Our results showed for the first time the bioavailability and potentiality of ChiNH/Q as a potent antimicrobial and anticancer agent against cancer cells. Our result provided evidence that ChiNH/Q could effectively reduce cellular DNMT expression levels and increase genomic global DNA methylation in HepG2 cancer cells. Our results suggest a potential clinical application of nanoparticles as antimicrobial and anticancer agents in combination cancer therapy.

Keywords: chitosan nanohydrogel, quercetin, cytotoxic activity, antimicrobial activity, DNA methylation, gene expression

Introduction

Human hepatocellular carcinoma (HCC) is the fifth leading common cancer in the world and it is the third main cause of cancer-related deaths, worldwide.^{[1](#page-10-0),2} In recent years, the role of environmental factors, including flavonoids, has drawn much atten-tion in the treatment of cancer.^{[3,](#page-10-2)[4](#page-10-3)} Quercetin $(3, 3', 4', 5, 7$ -pentahydroxyflavone) (Q) is one of the most important natural dietary polyphenolic flavonoids which is abundantly

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found in fruits and vegetables. It has been reported that Q could exert distinct antioxidant, anti-inflammatory, and anti-cancer effects and its role in preventing cancer has been addressed previously.^{[5](#page-10-4)-8} Moreover, it is well documented that Q is a potent bioflavonoid that exerts suppressing effects on the growth of several types of cancers including liver, esophagus, bladder, and colon cancer. [5](#page-10-4) However, the dualistic character of Q, regarding its role in the proliferation of cancer cell lines, has been investigated, which may affect present views on Q supposed beneficial anti-cancer effect. From a mechanistic viewpoint, it has been suggested that Q may affect the carcinogenic potential of various compounds. Many studies have shown that Q activates the metabolism of polyaromatic hydrocarbons and estrogens, which leads to the formation of catechol and subsequent oxidation metabolites known to be involved in carcinogenesis. $9-11$ Q already possesses a catechol group, which, due to its oxidant properties, provides the basis for the oxidizing toxic effects of this flavonoid.^{[12](#page-10-8)} Moreover, many studies pointed at possible adverse health effects caused by Q and reported that Q can be genotoxic and also play a mutagenic role in various cultured cells.^{[13](#page-10-9)[,14](#page-11-0)} Previous studies have shown that Q is also a potent chemical agent that could cause apoptosis in various cancer cells and it showed high antimicrobial properties.[15](#page-11-1) Moreover, it has been suggested that the anti-cancer effects of polyphenols resulted from the impact of these flavonoids on epigenetic changes including DNA methylation.¹⁶ Alteration in DNA methylation plays an important role in the biology of cancer and it mainly inhibits the expression of tumor suppressor genes in cancer cells, including liver cancer. [17](#page-11-3)[–20](#page-11-4) DNA methylation is a chemical modification that is mainly generated in the promoter CpG islands of the genome by the action of a family of enzymes known as DNA methyltransferases (DNMTs), including DNMT1, DNMT3A, and DNMT3B.^{[21,](#page-11-5)22} DNMT1 is the most abundant DNMT which is involved in faithful transmission and the maintenance of DNA methylation in daughter cells during cell divisions. DNMT3A and DNMT3B act as the most potent global *de novo* methyltransferases that prefer to act on unmethylated CpG islands.^{[22](#page-11-6)–24} Numerous studies suggested that DNMTs are up-regulated to different levels in various cancers, including $HCC^{22,25}$ $HCC^{22,25}$ $HCC^{22,25}$ $HCC^{22,25}$ $HCC^{22,25}$ Moreover, many reports have demonstrated that alteration in regional and global genomic DNA methylation patterns occur in early stages in liver tumorigenesis and might associate with the pathogenesis of HCC. 26 In vivo and in vitro studies have demonstrated that polyphenols such as Q could alter hepato-cytes metabolism and DNMTs' activity.^{[27](#page-11-10),[28](#page-11-11)} Also, molecular

docking studies have demonstrated that Q modulates the activity of DNMTs and can function as a competitive inhibitor of the active site of DNMTs.²⁹ Moreover, it has been suggested that Q could induce apoptosis via re-expression and inhibition of DNA methylation of apoptotic genes in cancer cell lines.³⁰ However, many studies indicated that the application of Q has been limited by its low solubility, low bioavailability, and also it showed poor percutaneous and circulation absorption.^{7[,31,](#page-11-14)32} Therefore, to solve this problem, many researchers have attempted to improve the bioavailability forms of Q via chemical modifications, including Q-loaded liposomes and nanoparticles. $32-34$ $32-34$ Quagliariello et al fabricated hyaluronic acid-based Q nanohydrogel and highlighted the antitumor and antiinflammatory properties of the Q nanocarriers in the regulation of breast cancer cell proliferation.³⁵ Advantages of using nanoparticles including the delivery of drugs and genes into various specific sites of the body could result in longer shelf-life of the drug at the target site, and improvement of intracellular absorption, and further, the stability of drugs has also been discussed.^{[8](#page-10-5)[,36-](#page-11-18)[38](#page-11-19)} Furthermore, targeted drug delivery through nanoparticles provides a new therapeutic approach, in which a drug that has been administered through intravenous or oral routes acts effectively, with lower side effects in the patient.^{[39](#page-11-20)–[42](#page-11-21)} Nanohydrogels have shown different characteristics, making them an excellent vehicle for the delivery of therapeutic agents. The use of the natural polymer chitosan, as a hydrogel scaffold, is very important due to its polymer compatibility, low toxicity, and environmental degradability. [43](#page-11-22) Chitosan is a linear amino polysaccharide, derived from deacetylation of chitin and various studies have shown that chitosan hydrogels contribute to appropriate delivery of drugs to specific sites. $32,44-48$ $32,44-48$ $32,44-48$ Moreover, it has been suggested that chitosan-derived hydrogels are sensitive to enzymatic degradation and can easily degrade in nature as well as in human body.^{[44](#page-11-23)-[48](#page-11-24)} In the current study, a novel chitosan-based quercetin containing nanohydrogel (ChiNH/Q) was fabricated (Patent) and after physicochemical characterization, its biocompatibility, cytotoxicity, antimicrobial and epigenetic aspects as a potential antitumor agent were investigated using well-characterized HepG2 human hepatoma cells.

Materials and Methods

All chemicals and reagents were purchased from Merck (Darmstadt, Germany), Sigma Aldrich (Gillingham, UK) and Gibco Invitrogen (Paisley, UK).

Preparation of Chitosan-Nanohydrogel (ChiNH) and Chitosan-Based Quercetin-Containing Hydrogels (ChiNH/Q)

The ChiNH nanoparticles were prepared by the ionic gelation method. Briefly, 1% chitosan solution (0.12%, 0.25%, 0.5% and 1% of chitosan) was prepared in acetic acid (0.2% v/v, Merck, Darmstadt, Germany), with vigorous stirring for 12 hours at pH=4.7. An aqueous solution of 0.5% tripolyphosphate (TPP) (0.12%, 0.25%, 0.5% and 1% of TPP) was also prepared separately. Both solutions were refined using a filter paper to remove insoluble particles (0.45μ) pore size). Then, 5 mL of the TPP solution was added to 20 mL of chitosan solution dropwise, and stirred for an additional 10 minutes. To produce ChiNH/Q, 5 mg of Q (0.62, 1.25, 2.5, and 5 mg of Q) was added to chitosan before the addition of TPP. After 12 hours of continuous stirring, the obtained particles were lyophilized, freeze-dried, and kept at 4ºC until further use. An optimal amount of ingredients (chitosan, TPP, and Q) for hydrogel formation were used.

Encapsulation Efficiency

About 0.02 g of the freeze-dried hydrogel was poured into the test tube, and after addition of 4 mL of 2M hydrochloric acid, the resulting solution was left in an ultrasonic apparatus for 30 min. Finally, the dispersed content was refined with filter paper and its volume was increased up to 10 mL by addition of double-distilled water (ddH₂O). To evaluate the amount of Q loaded in the formulation, the sample's absorbance was measured, using the UV/VIS spectrophotometer at 254 nm, and calculated by the equation below.^{[48,](#page-11-24)[49](#page-11-25)}

Encapsulation Efficiency(%) = 100 $\chi \left(\frac{\text{mass of loaded quercetin}}{\text{mass of initial quercetin}} \right)$

Particle Physicochemical Characterization

Size, Polydispersity Index (PDI) and Zeta Potential Size distribution and zeta potential of the nanoparticles were studied using dynamic light scattering (DLS) and a Malvern Zetasizer Nano-range instrument (Malvern Instruments Ltd., Malvern, UK), respectively. The colloidal emulsions were diluted in ddH₂O (1:1000, v/v) and studied in triplicate at 25°C, at a fixed angle, using this instrument.

Scanning Electron Microscopy (SEM)

To conduct the SEM imaging, freeze-dried nanoparticles (left at −50°C for 24 h) were left on a stub, sputter-coated with gold, and examined at 15 kV with a 6300 field emission scanning electron microscope (Hitachi, S-4160).

Fourier-Transform Infrared (FTIR) Spectroscopy

The infrared absorption spectra of the samples including chitosan solution, TPP, ChiNH, and ChiNH/Q were studied using FTIR (BRUKER, TENSOR 27. Germany).

X-Ray Diffraction (XRD) Test

The pattern of the ChiNH/Q and the formulation of ChiNH underwent X-ray diffraction to record possible peaks of the nanogel. The samples were finely pulverized in a glass substrate, using UV-filtered (Ni-filtered Cu Ka radiation) in the range of 2 theta (2 θ) at 2–50 \degree C, and the experimental parameters were set as follows: Voltage of 40 kV; current of 20 mA; and angular velocity of 4°/min. The XRD patterns of Q, ChiNH, and ChiNH/Q were recorded using the Dmax 2100 Rigaku diffractometer (Rigaku Americas Co., TX, USA).

Release Kinetics

The amount of Q emission was determined, using a dialysis bag. To this end, the dialysis bag was immersed in ddH2O at 25°C for 24 h. About five mL of the prepared formulation solution (equivalent to 1.5 mg of Q) and donor compartment was left in the dialysis bag (12 KD). The loaded bag was soaked in 90 mL distilled water containing 30 mL ethanol as an acceptor compartment. The acceptor compartment agitated at 600 rpm. At specified intervals, 1 mL of the release medium was separated at time intervals of 15, 30, 45, 60, 90, 120, 150, 180, 210, 240, and 300 min for analysis and immediately replaced with 1 mL of the fresh medium ($pH = 7.4$). The graph for the Q levels was plotted against time. The mechanism of release of the Q from the formulation was investigated by the Korsmeyer–Peppas equation using the following formula:

where M_t is indicating the amount of Q at any given time, M_0 is the Q amount at the beginning of the test, t is the release time, *K* is the kinetic constant for the release, and the value of n is the characteristic power for the release. The release mechanism can be described based on the value of n and indicates the Fickian and non-Fickian (anomalous) behaviors. For spherical particles, $n \leq 0.43$ indicates a case I transport (Fickian release), while $n = 0.85$ indicates a case II transport (non-Fickian release). Intermediate values of n, ie, $0.43 \le n \le 0.85$, indicate a non-Fickian release and the release follows the combination of the mechanisms of diffusion and

inflation.^{[50](#page-11-26),51} The linear graph was plotted using the Ln $\frac{M_l}{M_0}$ at the final time against Ln(t). The cumulative amount of Q released from the formulation was calculated at different intervals by the following equation and the graph for the cumulative amount was plotted against time.

Cumulative release(
$$
\%
$$
) = $\sum_{t=0}^{t} \frac{Mt}{M0} \times 100$

pH Test

To evaluate pH changes, 0.5 g of formulations with and without Q was added to 10 mL ddH₂O, and the pH was determined throughout 0–90 (0, 1, 2, 7, 14, 30, 45, 60, and 90) days.

HepG2 Cell Culture and MTT Assay

Human hepatoma HepG2 cell line was purchased from the National cell bank of Iran (NCBI, Pasteur Institute, Tehran). Cells were cultured in RPMI 1640 medium supplemented with 10% FBS, 2mM Gln, penicillin (100 U/mL), and streptomycin (100 μg/mL) in humidified 5% CO2, the atmosphere at 37°C temperature. For ChiNH, Q, and ChiNH/Q treatment, cells were seeded into six-well plates at a density of 3×10^4 cells and allowed to attach for 24 h. The viability of HepG2 cells was determined by MTT test using 5 different concentrations of 10, 50, 100, 500 and 1000 µg/mL of ChiNH, Q and ChiNH/Q for 48h based on the method described by Rashidipour et al.⁵² All experiments were performed in triplicate and the data were expressed as mean \pm SD.

Antimicrobial Activity of ChiNH/Q Microbial Strains and Culture

To evaluate the antimicrobial effects of ChiNH/Q, five strains of the pathogen, including *Staphylococcus aureus* (*S. aureus*) (ATCC 12600), *Listeria monocytogenes* (*L. monocytogenes*) (PTCC 1297), *Escherichia coli (E. coli)* (PTCC 1395), *Klebsiella pneumoniae* (*K. pneumoniae*) (PTCC 1290) and *Candida albicans (C. albicans)* (PTCC 5072) were obtained from the National Cell Bank of Iran (NCBI, Pasteur Institute, Tehran).

Antimicrobial effects of broth microdilution were used according to the CLSI standard method.^{[53,](#page-12-1)54} Briefly, to examine the Minimum Inhibitory Concentration (MIC), ChiNH/Q and blank (1000 µg/mL), Q (5000 µg/mL), Ceftriaxone/ Fluconazole (200 µg/mL) sterile media, Mueller-Hinton broth for bacteria and Sabouraud Dextrose broth agar (SDA) for *C. albicans* were prepared. Then, different concentrations of stocks were added to the wells and the microbial suspension equivalent to Half McFarland $(1.5\times10^8$ CFU/mL) was added to the wells and kept in a 37°C incubator for 24 hours. After this time, salts of 2,3,5- Triphenyltetrazolium chloride were used to evaluate microbial viability. Pink wells indicate microbial life and non-color wells indicate microbial inhibition.^{[53](#page-12-1)} After calculating the MIC, 5 µL was cultured on Mueller-Hinton agar (for bacteria) and SDA (for *C. albicans*) with 24 min incubation, and then incubated for 24 h. At 37°C, the first concentration where no colony was found, reported as the Minimum Bactericidal Concentration and Minimum Fungicidal Concentration (MBC/MFC).

RNA Extraction and Quantitative Real-Time RT-PCR

Total RNA was extracted from HepG2 cells using the TriPure isolation reagent (Roche Applied Science, Germany) according to the manufacturer's instructions. The integrity of purified RNA was assessed on 2% formaldehyde containing 1.5% agarose, and purified RNA was stored at −80°C until use. The complementary DNAs (cDNAs) were synthesized from 2 µg total RNA by reverse transcriptase (RT) using oligodT primers and M-MuLV RT (MBI Fermentas, Lithuania) in a 20μL final reaction volumes according to the manufacturer's provider. Transcript levels of mRNA encoding DNMT1, DNMT3A, and DNMT3B were analyzed by quantitative real-time RT-PCR assay using a Corrbet sequence detection system (Rotor gene 6000). SYBR Green-based analysis and Master Mix (ABI, UK) were used and all reactions were carried out in triplicate. The primers information and conditions for DNMTs real-time RT-PCR procedures have been previously described.^{[22](#page-11-6)} The GAPDH was used as a reference gene and the relative gene expression levels were determined using the $2 - \Delta \Delta CT$ standard method.⁵⁵

Analysis of Global Genomic DNA Methylation

Global DNA methylation was quantified in DNA isolated from HepG2 cells using a 5-mC DNA ELISA kit (Zymo Research, Freiburg, Germany) according to the manufac-turer's instructions as previously described.^{[22](#page-11-6)}

Statistical Analysis

SPSS 21 analytic software (SPSS, Inc., Chicago) and Graph Pad Prism (Version 8.01) were used for data

analysis. All data were presented as mean \pm standard deviation (SD) from three independent experiments. Data were analyzed using one-way ANOVA followed by Tukey's multiple comparison tests. Differences with p-value ≤0.05 were set as the level of significance.

Results and Discussion

We hypothesized that "a novel quercetin Q nanohydrogel, using Chitosan (ChiNH/Q) would have more biological properties for pathogenic microbial strains and cancer cells than free Q". Many studies effectively highlight the high antimicrobial properties of Q and ability of Q to modulate epigenetic modifications particularly DNA methylation and restore gene expression. However, the application of Q is limited by several factors including low bioavailability, low solubility, and low circulation absorption, and also application forms of Q. Numerous methods including nanoliposomal and nanoparticle-based targeted delivery to tumors, are actively being sought to develop the bioavailability of Q. To the best of our knowledge, no previous studies have addressed the bioavailability, cytotoxicity, antimicrobial properties, and effect of Q nanohydrogels on DNA methylation in HCC.

Therefore, our study is the first report for the synthesis of ChiNH/Q. Due to the many important characteristics of chitosan, including cationic characters and primary amino groups, which are responsible for many associated properties such as transfection and controlled drug release, it has been suggested that chitosan is the most important poly-saccharide for various drug delivery purposes.^{[56](#page-12-4)} The SEM image of the freeze-dried hydrogel sample was prepared which indicated that ChiNH was successfully synthesized from chitosan ([Figure 1A](#page-4-0)). The compact and dense matrix could be due to the formation of electrostatic bondings between the TPP functional groups and the amino groups in the chitosan. The hydrodynamic size, zeta potential, and PDI of the produced particles were measured [\(Table 1\)](#page-5-0). The encapsulation efficiency was found to be 62.24% and the percentage of Q in chitosan-based quercetin-containing hydrogels was 1.42%. The size of the ChiNH/Q was 743.6 nm and the size of ChiNH was 992.7 nm. The changes in the PDI, in the ChiNH/Q and ChiNH were 0.507 and 0.537, respectively. These changes indicate a uniform dispersion of particles in the formulation of ChiNH/Q. $57-59$ $57-59$ $57-59$ Our results demonstrated that ChiNHs and loading Q could affect the particle sizes so that ChiNHs showed

Figure 1 (**A**) The scanning electron microscopy (SEM) micrographs at 15 kV of unloaded ChiNH and ChiNH/Q. (**B**) The FTIR spectrum analysis of ChiNH/Q, ChiNH, Q, TPP and chitosan. (**C**) The XRD peaks of ChiNH and ChiNH/Q formulation. **Abbreviations:** ChiNH, chitosan-based nanohydrogel; Q, quercetin; ChiNH/Q, chitosan-based quercetin nanohydrogel.

	0 day		60 days	
	ChiNH/Q	ChiNH	ChiNH/Q	ChiNH
Diameter (nm)	743.6	992.7	912	876.4
Polydispersity index (PDI) Zeta potential (mV)	0.507	0.537	0.348	0.649
	12.1	3.7		

Table 1 The Physicochemical Characteristics of ChiNH/Q and ChiNH as a Function of Time (0 and 60 days)

Abbreviations: nm, nanometer; mV, millivolt; ChiNH, chitosan-based nanohydrogel; ChiNH/Q, chitosan-based quercetin nanohydrogel.

the largest size (992.7 nm), whereas, ChiNHs/Q showed the smallest size (743.6 nm) [\(Table 1](#page-5-0)). The smaller size of ChiNHs/Q might be due to the crosslinking between the sodium-TPP functional group, the chitosan amine groups, and the hydroxyl functional groups in the Q. For practical purposes, the physical characteristics of nanoparticles including size, shape, structure, and coating have significant effects on the cytotoxicity, distribution, and also cellular uptake of nanoparticles. It has been suggested that the nanoparticles of comparably smaller sizes are better because of the ratio of surface to volume as well as colloidal stability increases as the particle size decreases.[57](#page-12-5) Zeta potential was determined as 12.1 mV and 3.7 mV for ChiNH/Q and ChiNH, respectively, given the presence of the anionic TPP compounds in nanohydrogels.

To investigate the chemical composition of the produced ChiNH in the presence and absence of Q, the FTIRs were separately studied [\(Figure 1B\)](#page-4-0). FTIR analysis indicated that the formation of bioactive groups surrounded the Q in the hydrogel cavities. The FTIR spectrum of ChiNH/Q presented 6 peaks especially at 1153–1415 cm⁻¹, 1563 cm⁻¹, 1638–1640 cm−1, 2925 cm−1 and 3237–3562 cm−1 for ChiNH/Q [\(Figure 1B](#page-4-0)). The broad and strong absorbance peaks of 1163–1216 cm⁻¹ indicate the presence of symmetrical and asymmetric stretching vibration, PO_3 group and P=O groups related to chitosan and sodium TPP, respectively.⁶⁰ The absorption peak at 3570 cm⁻¹ was assigned to the existing N-H stretching of amid groups of chitosan. The signals observed at 1617 cm^{-1} attributed to amid II (N-H bonding) of chitosan. Absorbance peaks at the wavelengths of 1638–1640 cm⁻¹ and 2925–3237 cm⁻¹ indicated the presence of symmetric and asymmetric tensile vibrational groups of $C=O$; 1455 cm⁻¹ was assigned to the aromatic C=C stretching and 1137 cm^{-1} indicated phenolic C-O stretching related to Q. Moreover, in the ChiNH/Q there was a strong absorbance peak at 627 cm^{-1}

corresponding to tensile vibrational of $NH₂$ -groups related to Q, and indicating that Q was included in the formulation, and the presence of this peak in the respective spectrum indicating cross-linking created by Q [\(Figure 1B\)](#page-4-0). These data demonstrate the formation of a chitosan-based quercetin nanohydrogel. Also, based on the FTIR data, the dispersion was potentially lower due to the interaction and loading of the Q inside the formulation cavities.

According to the results of XRD graphs, no specific peak was observed on the patterns of the ChiNH and ChiNH/Q formulations [\(Figure 1C\)](#page-4-0). As shown in [Figure 1C](#page-4-0), this observation explains that Q is encapsulated or dispersed among the chitosan-sodium TPP polymers which are linked by an interconnected ionic network, and an amorphous complex is formed through intermolecular interaction in the matrix. Various factors affect the release of nanoparticles, including the particle size, molecular weight of the polymer, and the interaction between the particles and the polymer. [60,](#page-12-7)[61](#page-12-8) To investigate the releasing mechanism, the final formulation was prepared under optimal conditions, and Q release was calculated at 15, 30, 45, 60, 90, 120, 150, 180, 210, 240, and 300-min intervals using the dialysis bag [\(Figure 2A\)](#page-6-0). In this test, at specified intervals, 1 mL of the receiving solution was separated for analysis and it immediately replaced with 1 mL of the fresh receiving solution (pH=7.4). The specific concentration of ChiNH/Q was poured into the dialysis bag (12 KD) and left in the container. The value of n was calculated, using the graphed equation of the line. According to [Figure 2A](#page-6-0) and the obtained data, 77.28% of the initial Q was released after 300 minutes. The Korsmeyer–Peppas mathematical model was used to calculate n, which was ultimately calculated at 0.74 ([Figure 2B\)](#page-6-0). This indicates that Q release from chitosan-based polymer follows the combination of two processes: penetration and erosion. The constant of release (K) was 1.61 and the correlation coefficient was 0.96 [\(Figure 2B](#page-6-0)).^{[52](#page-12-0)} As a test of stability, pH showed that the H^+ concentration in the formulation increased by approximately 1.05 times from day 0 to day 90, which could be due to the presence of chitosan hydrolysis and release of the glucosamine and oligosaccharides [\(Figure 3](#page-6-1)).

Our study indicated for the first time, the cytotoxicity effect of encapsulated Q by ChiNHs against HepG2 cells using the MTT assay. The optical microscopy was employed to examine the influences of ChiNH, Q and ChiNH/Q on the morphological changes of HepG2 cells [\(Figure 4](#page-7-0)). Our data indicated that ChiNH/Q at concentrations of 50, 100, 500, and 1000 µg/mL significantly

Figure 2 The kinetic release of Q from ChiNH in PBS. (**A**) The cumulative release of Q in a 300-min time course was calculated as a percentage of total Q loaded into chitosan nanohyrogel. (**B**) the Korsmeyer–Peppas mathematical model was used to calculate n.

decreased the HepG2 cell viability in a dose-dependent way, as compared with untreated control cells $(P<0.05)$ [\(Figure 5](#page-7-1)). Furthermore, we found that ChiNH/Q significantly increased drug cytotoxicity on HepG2 cells, as compared with free Q (p<0.05) [\(Figure 5](#page-7-1)). Moreover, the IC_{50} values of ChiNH/Q against HepG2 cells were significantly lower than free Q at 50 μ g/mL (p<0.05) [\(Figure 5\)](#page-7-1). Then, 100 µg/mL (lethal-dose) of each of ChiNH/Q, and

Figure 3 The pH of ChiNH and ChiNH/Q formulation throughout 90 days. **Abbreviations:** ChiNH, chitosan-based nanohydrogel; ChiNH/Q, chitosan-based quercetin nanohydrogel.

free Q showed 50.7% and 57% cell viability at 48 hours, respectively. Whereas, 50 µg/mL (sub-lethal dose) of each of ChiNH/Q and free Q showed 65% and 76.5% cell viability at 48 h, respectively. These data have proved the efficiency of ChiNH/Q as an anticancer agent with low IC_{50} against HepG2 cells. The mechanisms by which nanoparticles can stop the growth of cancer cells are still not clear. However, experimental studies indicated that Q-nanoparticles induced antioxidant activity and more efficient interaction of Q with proteins. 62 Moreover, in vitro studies has been indicated that Q-nanoparticles are a promising candidate for the delivery of the anticancer drug delivery in cancer cells and show impressive cyto-toxicity for cancer cells.^{[63](#page-12-10)}

Our study for the first time evaluated the antimicrobial activity of ChiNH/Q against different pathogenic bacterial strains, including *K. pneumoniae* PTCC 1290, *E. coli* PTCC 1395, *S. aureus* ATCC 12600, *L. monocytogenes* PTCC 1297 and the yeast *C. albicans* PTCC 5072, using broth microdilution assay and according to the CLSI stan-dard method.^{[53](#page-12-1)} According to the results in [Table 2](#page-8-0), the

Figure 4 The optical microscopy images of the morphological changes of HepG2 cells after treatment with 50 µg/mL of ChiNH, Q and ChiNH/Q for 48 h.

antibacterial and antifungal results showed that ChiNH/Q suppressed the growth of Gram-positive bacterium *L. monocytogenes* and *C. albicans* to greater extents compared to free Q as evidenced by MIC 250 and 1000, respectively, while free Q resulted in MIC equal to 1250 in both of *L. monocytogenes* and *C. albicans* [\(Table 2\)](#page-8-0). These results are likely to be due to the proper enclosure of Q in the nanohydrogel network, and the better effects of encapsulated Q could be attributed to its slower release in the polymer matrix.^{[64](#page-12-11)} Moreover, our results indicated that for two Gram-negative bacterial strains, including *E. coli* and *K. pneumoniae*, the MIC of free Q was 312.5 and 156.25, respectively. These results demonstrated that free Q showed a greater inhibitory effect than ChiNH/Q on these Gram-negative bacteria ([Table 2](#page-8-0)). Also, for Grampositive bacteria like *S. aureus* the MIC was >1000 after treating with free Q and ChiNH/Q. These data showed that *S.aureus* is resistant to all treatments, possibly due to the complexity of the cell wall structure of this bacterium and many studies have shown that Gram-positive bacteria are more resistant to the penetration of nanoparticles. $65-68$ $65-68$ Since, Gram-positive bacteria such as *S. aureus* have

Figure 5 The cytotoxicity of ChiNH (nano-blank), Q and ChiNH/Q in HepG2 cancer cells. ChiNH/Q (combination of ChiNH and Q) synergistically improved cytotoxicity in HepG2 cells line in a dose-dependent manner at 48 h. All data repeated in triplicate shown as means ± SD.

Abbreviations: ChiNH, chitosan-based nanohydrogel; Q, quercetin; ChiNH/Q, chitosan-based quercetin nanohydrogel.

a much thicker layer of peptidoglycan than Gramnegative bacteria, including *E.coli*, the penetration of nanoparticles into the wall of Gram-negative bacteria is easier due to the lower thickness.^{[69](#page-12-14)} Our data indicated that ChiNH/Q showed a potent antimicrobial effect against *L. monocytogenes* and C.albicans. Moreover, the results are interesting since the values of MIC caused by ChiNH/ Q are more close to Ceftriaxone/Fluconazole, as positive control for *L. monocytogenes* and *C. albicans* compared to free Q [\(Table 2](#page-8-0)). Our results are in line with Milanezi et al that Q-capped gold nanoparticles demonstrated a higher antimicrobial activity than free Q in pathogenic strains.^{[70](#page-12-15)} It has been suggested that the small size of the nanoparticles makes them suitable targets for antimicrobial and intracellular targeting purposes.^{[71](#page-12-16)} Moreover, nanogels and microgels as drug carriers are capable of carrying significant amounts of pharmaceutical compounds and protecting them against proteolytic degradation.^{[72](#page-12-17)} Overall, our data indicated that ChiNH/Q might be a potent antimicrobial agent with a strong inhibitory effect on pathogenic microorganisms and could be used as a potential alternative antibiotic in medical and therapeutic applications.

We evaluated for the first time the application of encapsulated Q by ChiNHs on the DNMTs' gene expression in HepG2 cells. To verify if the DNMTs'expression is influenced by ChiNH, ChiNH/Q, and free Q treatment, we selected a sub-lethal dose of 50 μ g/mL and HepG2 cells were incubated with 50 µg/mL of ChiNH/Q, ChiNH, and free Q for 48 h. We observed variable effects of ChiNH, ChiNH/Q, and free Q on all the three DNMT´s gene expressions (DNMT1, DNMT3Aand DNMT3B) in HepG2 cells. As shown in [Figure 6A](#page-8-1) and [Table 3,](#page-9-0) our study showed that treatment of HepG2 cells with ChiNH/ Q resulted in a decrease in mRNA transcription levels of DNMT1, DNMT3A, and DNMT3B by 68%, 59%, and 65%, respectively, as compared to control cells $(P<0.05)$

Microorganisms Conditions	K. pneumoniae PTCC 1290 MIC MBC	E. coli PTCC 1395 MIC MBC	S. aureus ATCC 12600 MIC MBC	L. monocytogenes PTCC 1297 MIC MBC	C. albicans PTCC 5072 MIC MFC
Blank	>1000 > 1000	1000>1000	>1000 > 1000	500500	1000 > 1000
Q	156.25312.5	312.5625	>5000 > 5000	1250 2500	1250 2500
ChiNH/O	5001000	1000 > 1000	>1000 > 1000	250 1000	10001000
Ceftriaxone/Fluconazole	1.56 1.56	3.126.24	12.550	12.550	100 200

Table 2 The Antimicrobial Effects of Quercetin-Chitosan Nanohydrogel

Abbreviations: PTCC, Persian Type Culture Collection; ATCC, American Type Culture Collection; MIC, minimum inhibitory concentration; MBC/MFC, minimum bactericidal concentration/minimum fungicidal concentration.

[\(Table 3](#page-9-0)). Moreover, ChiNH/Q showed significant effects on the DNMT1 expression level in HepG2 cells, compared to ChiNH-treated cells (P<0.05) ([Figure 6A\)](#page-8-1). According to previous reports, the DNMT1 expression is a key element in liver malignancy and significantly correlated with the poor prognosis of HCC.⁷³ Furthermore, experimental studies suggested an indispensable role of DNMT1 as a key genomic regulator of liver growth and regeneration.⁷⁴ Moreover, the statistical analysis of our study showed that Q similar to ChiNH/Q could suppress the expression

level of DNMT3A (70%-reduced) and DNMT3B (63% reduced) in HepG2 cells compared to control cells (P<0.05) [\(Figure 5A\)](#page-7-1) [\(Table 3\)](#page-9-0). Interestingly, our results indicated that the average expression level of all three DNMTs (T1/3A/3B) was effectively reduced by ChiNH/ Q (65% reduced) compared to control cells (P<0.05) [\(Figure 6B\)](#page-8-1) [\(Table 3](#page-9-0)). We observed a trend for the coordinated pattern of all three DNMTs' average inhibition expression, using ChiNH/Q, in comparison to free Q ([Figure 6B](#page-8-1)). As expected, our results demonstrated

Figure 6 (**A**) The relative expression of DNMT1, DNMT3A and DNMT3B in HepG2 cells treated with ChiNH, Q and ChiNH/Q, measured by real time-PCR. The expression of each gene was normalized to GAPDH. Cells treated with only RPMI 1640 were considered as controls and the rate of expression of the genes of interest in other groups was calculated. The data were presented as mean ± SD and each experiment was conducted in triplicate. The results were analyzed by one-way ANOVA and Tukey's post hoc test. The significance level is considered as p <0.05. The bars marked with different letters (a, b, c) are significantly different from other samples. (**B**) A summary of the change in the average expression of all 3 DNMTs (DNMT1, DNMT3A and DNMT3B) in HepG2 cells treated with ChiNH, Q and ChiNH/Q for 48 h. **Abbreviations:** C, control; ChiNH, chitosan-based nanohydrogel (nano-blank); Q, quercetin; ChiNH/Q, chitosan-based quercetin nanohydrogel.

Gene	ChiNH		ChiNH/Q
DNMTI	1.137 ± 0.4381^a	1.146 ± 0.7275^a	0.3239 ± 0.1011 ^b
DNMT3A	0.5819 ± 0.3636^{ab}	0.3057 ± 0.1833^b	$0.4104 \pm 0.3070^{\circ}$
DNMT3B	0.7084 \pm 0.2053 $^{\rm ab}$	0.3713 ± 0.3193^b	0.3485 ± 0.0482^b
T1/3A/3B	0.7973 ± 0.4095^{ab}	0.6121 ± 0.5902 ^{bc}	0.3562 ± 0.1724 ^c

Table 3 The Effects of ChiNH, Q and ChiNH/Q on DNMTs mRNA Gene Expression Levels in the HepG2 Cancer Cells. The Values are Represented as Mean ± SD

Note: a,b,cDifferent letters are significantly different from other samples at p <0.05.

Abbreviations: C, control; ChiNH, chitosan-based nanohydrogel (nano-blank); Q, quercetin; ChiNH/Q, chitosan-based quercetin nanohydrogel.

that the decreased DNMTs' expression level was observed only in ChiNH/Q-treated HepG2 cells compared to ChiNH-treated cells ([Figure 6B](#page-8-1)). Overall, based on these results, downregulation of DNMTs was effectively observed in the ChiNH/Q-treated HepG2 cells, and ChiNH/Q exhibits a greater effect on the reduction of the expression level of DNMTs while compared to free Q. Many studies have reported the increased DNMTs' expression levels in HCC, while reducing the expression of DNMTs by ChiNH/Q could help to control HCC.[75](#page-12-20)

This report demonstrated for the first time, the application of ChiNH/Q on the global genomic DNA methylation profile in HepG2 cells. Our results indicated that overall about 0.995% of the cytosine residues were methylated in the genome of the media-only treated (control) HepG2 cells ([Figure 7](#page-9-1)). Moreover, we found that the level of methylated cytosine in ChiNH/Q-treated cells was

Figure 7 The Effects of 50 µg/mL concentration of ChiNH, Q and ChiNH/Q with concentrations on global DNA methylation in HepG2 cells. Values represent mean ± SD of three experiments. The bars marked with asterisk are significantly different as verified by Tukey's honestly significant difference multiple comparison test (p < 0.05).

Abbreviations: C, control; ChiNH, chitosan-based nanohydrogel (nano-blank); Q, quercetin; ChiNH/Q, chitosan-based quercetin nanohydrogel.

significantly higher (1.01%) than those with Q and ChiNH-treated cells (0.993% and 0.992%, respectively) (p<0.05) ([Figure 7](#page-9-1)). As expected, these results indicated that ChiNH/Q effectively increased the global DNA methylation of HepG2 cells, compared to ChiNH and free Q-treated cells. These results suggested that ChiNH/ Q always displayed further efficacy, in comparison to free Q in decreasing DNMTs' gene expression and also increasing the global DNA methylation in HepG2 cancer cells. Many studies also demonstrated that Q can exert anti-cancer effect by regulating epigenetic modifications including, miRNA expression and DNA methylation, and enhance the sensitivity of tumor cells to chemotherapy.^{[76,](#page-12-21)[77](#page-12-22)} Other studies also reported the antiinflammatory properties of Q nanoparticles and demonstrate that biodegradable Q nanoparticles increases the cellular uptake of chemotherapy drugs into cancer cells and enhances their cytotoxicity and antitumor properties.^{[78](#page-12-23)} Global DNA hypomethylation is a common epigenetic change in HCC and plays an important role in elevating chromosomal instability, and initiation and progression of HCC.^{[26](#page-11-9),79} Overall, our study successfully explains that ChiNH/Q always shows a higher efficacy than free Q, and ChiNH/Q can effectively reduce cellular DNMTs' expression level and also increase genomic global DNA methylation in HepG2 cancer cells. By linking polyphenol nanoparticles and DNA methylation, our current study provides some insight into the capacity of nanoparticles in the delivery of anti-tumor agents regarding epigenetic modification of HCC and also can open new insights to clinical potential of nanoparticles in targeted-therapy in cancer.

Conclusions

In conclusion, the formulation of novel Q nanohydrogel using chitosan (ChiNH/Q) detailed in this study was assessed by PDI, DLS, SEM, FTIR, and XRD. Encapsulation efficiency was confirmed by UV/VIS

spectrophotometer. The physicochemical characterization of ChiNH/Q showed that the size of the ChiNH/Q was 743.6 nm. The FTIR peak showed that the presence of carboxyl, $P=O$, $C=O$, and $NH₂$ groups were important factors in the synthesis of ChiNH/Q, and XRD data detected the encapsulation of Q in the ChiNH. The ChiNH/Q produced in this study exerted significant cytotoxic effects against HepG2 cancer cells with IC_{50} values of 100 μ g/mL. Also, ChiNH/ Q applied potent antimicrobial effects against Gram-positive bacteria and *C. albicans*, and it might be a promising antimicrobial agent in medical applications. Moreover, our results showed that ChiNH/Q increased availability of Q and exhibited a greater effect on the reduction of DNMTs' gene expression and also ChiNH/Q increased in global DNA methylation in HepG2 cancer cells. Our study provides new insights into the epigenetic mechanisms by which nanohydrogels containing flavonoids, effectively influence the gene expression in HCC. Also, our data indicated that the synthesis of Q nanoparticles was efficient and could be considered as an agent carrier in targeted therapy in liver cancer. However, further investigations including in vitro and in vivo models are needed to evaluate the potential application of the ChiNH/Q in nano-therapy approaches.

Abbreviations

Q, quercetin; ChiNH, chitosan-nanohydogel; ChiNH/Q, chitosan-based quercetin nanohydrogel; PCR, polymerase chain reaction; MIC, minimum inhibitory concentration; SDA, Sabouraud dextrose broth agar; MBC, minimum bactericidal concentration; MFC, minimum fungicidal concentration; SEM, scanning electron microscope; DLS, dynamic light scattering; FTIR, Fourier-transform infrared spectroscopy; XRD, x-ray diffraction; UV/VIS, ultraviolet–visible spectroscopy; ELISA, enzyme-linked immunosorbent assay; DNMTs, DNA methyltransferases; DNMT1; DNA methyltransferase 1; DNMT3A, DNA methyltransferase 3A; DNMT3B, DNA methyltransferase 3B; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; miRNA, microRNA.

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Author Contributions

All authors made a significant contribution to the work reported, whether that is in the conception, study design, execution, acquisition of data, analysis and interpretation, or in all these areas; took part in drafting, revising or critically reviewing the article; gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

Disclosure

The authors declare that there is no conflict of interest.

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