Dendrosomal curcumin nanoformulation downregulates pluripotency genes via miR-145 activation in U87MG glioblastoma cells

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Abstract: Glioblastoma is an invasive tumor of the central nervous system. Tumor recurrence resulting from ineffective current treatments, mainly due to the blood–brain barrier, highlights the need for innovative therapeutic alternatives. The recent availability of nanotechnology represents a novel targeted strategy in cancer therapy. Natural products have received considerable attention for cancer therapy because of general lower side effects. Curcumin is a new candidate for anticancer treatment, but its low bioavailability and water solubility represent the main disadvantages of its use. Here, curcumin was efficiently encapsulated in a nontoxic nanocarrier, termed dendrosome, to overcome these problems. Dendrosomal curcumin was prepared as 142 nm spherical structures with constant physical and chemical stability. The inhibitory role of dendrosomal curcumin on the proliferation of U87MG cells, a cellular model of glioblastoma, was evaluated by considering master genes of pluripotency and regulatory miRNA (microribonucleic acid). Methylthiazol tetrazolium assay and flow cytometry were used to detect the antiproliferative effects of dendrosomal curcumin. Annexin-V-FLUOS and caspase assay were used to quantify apoptosis. Real-time polymerase chain reaction was used to analyze the expression of OCT4 (octamer binding protein 4) gene variants (OCT4A, OCT4B, and OCT4B1), SOX-2 (SRY [sex determining region Y]-box 2), Nanog, and miR-145. Dendrosomal curcumin efficiently suppresses U87MG cells growth with no cytotoxicity related to dendrosome. Additionally, the accumulation of cells in the SubG1 phase was observed in a time- and dose-dependent manner as well as higher rates of apoptosis after dendrosomal curcumin treatment. Conversely, nonneoplastic cells were not affected by this formulation. Dendrosomal curcumin significantly decreased the relative expression of OCT4A, OCT4B1, SOX-2, and Nanog along with noticeable overexpression of miR-145 as the upstream regulator. This suggests that dendrosomal curcumin reduces the proliferation of U87MG cells through the downregulation of OCT4 (octamer binding protein 4) variants and SOX-2 (SRY [sex determining region Y]-box 2) in an miR-145-dependent manner.

Keywords: glioblastoma, pluripotency gene, miRNA, dendrosomal curcumin

Introduction
Glioblastoma is characterized by a high potential for growth and invasion. Insensitivity of this tumor to general therapies, including radio- and chemotherapy, leads to a high rate of recurrence. Until now, little was known about the genetic mechanisms involved in gliomagenesis.1–4 A positive association between the overexpression of stem cell marker genes and progression/metastasis of tumors triggered the idea that targeting of self-renewal pathways can be applied as a powerful approach in cancer treatment.5

Transcription factors, OCT4 (octamer binding protein 4), SOX-2 (SRY [sex determining region Y]-box 2), and Nanog play crucial roles in the maintenance of stemness in both embryonic and somatic stem cells.6 Several studies have reported the oncogenic...
role of these transcription factors in glioblastoma cells.\textsuperscript{2,3,7,8} Regulation of these pluripotency factors is under the control of a subset of micro ribonucleic acids (miRNAs), which are favorably expressed in undifferentiated stem cells. The promoter regions of these miRNAs and pluripotency factors are implicated in a precise reciprocal interaction giving rise to cell pluripotency or differentiation.\textsuperscript{9} \textit{miR-145} is well understood in stemness pathways. It has been demonstrated that \textit{miR-145} induces differentiation in stem cells by suppressing the promoter of \textit{OCT4} and \textit{SOX-2}.\textsuperscript{9,10} This explains the cause of low level \textit{miR-145} expression in most cancers.\textsuperscript{11} There is a considerable attention to replacing the chemical anticancer treatment agents with plant-based formulations to mitigate the undesirable effects during the prolonged treatment of patients. Curcumin is a hydrophobic compound extracted from the rhizome of \textit{Curcuma longa}, which is employed as a spice and coloring agent in most Asian countries. A growing body of evidence has substantiated the antitumor properties of curcumin in various cancer cells through interaction with multiple signaling pathways.\textsuperscript{12–15} However, low water solubility, poor uptake and tissue distribution remain major obstacles, limiting the use of curcumin as an anticancer treatment agent.\textsuperscript{16} Recently, scientists have revealed several strategies, such as loading synthetic analogs from turmeric, designing metabolic inhibitors and liposomal formulations, and nanoparticles of curcumin, to overcome these problems.\textsuperscript{17–20} Dendrosome was introduced as a novel neutral, amphipathic, and biodegradable nanocarrier for a gene delivery system in the authors’ research group.\textsuperscript{21,22} The high potential of dendrosome as a gene porter led to the hypothesis that it can be exploited as a vehicle for curcumin delivery. The anticancer properties of dendrosomal curcumin (DNC) have been explored in mice models of fibrosarcoma and colon cancer.\textsuperscript{21–24} It was demonstrated that dendrosome improves the solubility of curcumin and facilitates its uptake by cancer cells, suggesting that this polymer could open new avenues for successful drug delivery in cancer therapy.\textsuperscript{21,24} The present study reports for the first time a full characterization of DNC and describes the underlying mechanism through which curcumin targets the pluripotency pathways of glioblastoma cancer cells. U87MG – a cellular model of grade IV glioblastoma – was used in this regard. The probable cytotoxicity of this formulation on normal stem cells and fibroblast cells was also evaluated to determine if DNC can be considered a safe preparation for use in clinical and pharmaceutical approaches.

**Material and methods**

**Chemicals and standards**

Curcumin was purchased from Merck KGaA (Darmstadt, Germany) with purity of 95%. Materials for the preparation of dendrosome were oleoyl chloride and polyethylene glycol 400 that was purchased from Sigma-Aldrich (St Louis, MO, USA), and chloroform and triethyl amine were purchased from Merck. All the solvents used were high-performance liquid chromatography (HPLC) grade from Merck; 85% formic acid was provided by Carlo Erba Reagents (Cornaredo, Italy). Water was purified by a Milli-Q® Plus system from EMD Millipore (Billerica, MA, USA). Phosphate-buffered saline (PBS) solution was from Sigma-Aldrich. Curcumin reference standard (10 mg) was provided by Extrasynthese (Genay Cedex, France).

**Polymeric micelle OA400 carrier (dendrosome) preparation**

**Synthesis of dendrosome**

The OA400 carrier was synthesized by esterification of oleoyl chloride (0.01 mol) and polyethylene glycol 400 (0.01 mol) in the presence of triethyl amine (0.012 mol) and chloroform as the solvent at 25°C for 4 hours. Triethylamine hydrochloride salt was filtered from its organic phase. Then, chloroform was evaporated from OA400 in vacuum oven at 40°C for 4 hours (Figure 1). Fourier transform infrared spectroscopy (Spectrum One, PerkinElmer, Inc., Shelton, CT, USA) analysis was performed to confirm the chemical structure of the synthetic OA400 carrier. The OA400 carrier was synthesized by esterification of oleoyl chloride and polyethylene glycol 400 with triethyl amine and chloroform as the solvent at 25°C for 4 hours. Triethylamine hydrochloride was filtered from its organic phase. Then, chloroform was evaporated from OA400 in vacuum oven at 40°C for 4 hours (Figure 1). Fourier transform infrared spectroscopy (Spectrum One, PerkinElmer, Inc., Shelton, CT, USA) analysis was performed to confirm the chemical structure of the synthetic OA400 carrier.

**Figure 1** Synthesis of the OA400 carrier.

**Abbreviation:** PEG, polyethylene glycol.

\[\text{O} \quad \text{H}_3\text{C} \quad -\text{(CH}_2\text{)}_7\text{C} \quad \text{-C} \quad \text{H} \quad \text{Cl} \quad + \quad \text{H} \quad -\text{(OCH}_2\text{CH}_2\text{)}_n\text{OH} \quad \text{PEG 400} \]

\[\text{OA400 carrier} \]

\[\text{H}_3\text{C} \quad -\text{(CH}_2\text{)}_7\text{C} \quad \text{-C} \quad \text{H} \quad \text{Cl} \quad + \quad \text{H} \quad -\text{(OCH}_2\text{CH}_2\text{)}_n\text{OH} \quad \text{PEG 400} \]

\[\text{OA400 carrier} \]
performed to confirm the dendrosomal chemical structure. The Fourier transform infrared spectrum of OA400 shows a stretching band of C–H aliphatic at 2,921 cm$^{-1}$, a stretching band of C–H unsaturated aliphatic at 3,002 cm$^{-1}$, and a stretching band carbonyl group of ester bands at 1,732 cm$^{-1}$. The C–H bending vibration of CH$_2$ and the C–H bending vibration of CH$_3$ can be seen at 1,462 cm$^{-1}$ and 1,350 cm$^{-1}$, respectively. The C–O stretching vibration was at 1,111 cm$^{-1}$ as a broad band. The C=C stretching vibration can be seen at 1,651 cm$^{-1}$ (Figure S1).

Preparation of the DNC

DNC preparation was performed using optimized protocol in the authors’ lab. Briefly, different weight/weight ratios of dendrosoic/curcumin ranging from 50:1 to 10:1 were examined before settling on a suitable ratio of 25:1. Curcumin was dissolved in various amounts of dendrosome and checked for absorbance spectra by ultraviolet spectrophotometry (Infinite® 200 PRO, Tecan, Mannedorf, Switzerland). Then, the appropriate mixture of dendrosome and curcumin was evaluated for excitation/emission value in comparison with curcumin dissolved in PBS and 1% methanol as the control samples. The loading of dendrosome nanocarriers with curcumin molecules was performed using Gou et al’s protocol. Briefly, curcumin and dendrosome were co-dissolved in 5 mL acetone; this solution was added into 5 mL PBS while stirring constantly. Then, the acetone was evaporated in a rotary evaporator. The curcumin/dendrosome micelle solution was sterilized using a 0.22 µm syringe filter (EMD Millipore). Finally, the prepared DNC were stored at 4°C in a light-protected condition until used. For in vitro experiments, DNC was diluted in complete culture medium.

In vitro characterization

Particle characterization of the dendrosomal formulation

DNC particle size was measured by dynamic light scattering (DLS) (Zetasizer® Nano ZS90; Malvern Instruments, Malvern, UK). Time correlation functions were analyzed to obtain the hydrodynamic diameter of the particles and the particle size distribution (polydispersity index [PDI]) using ALV-60X0 software V.3.X (Malvern). Autocorrelation functions were analyzed by the cumulant method fitting a single exponential to the correlation function to obtain the mean size, PDI, and distribution method (to fit a multiple exponential to the correlation function to obtain particle size distributions). Scattering was measured in an optical quality 4 mL borosilicate cell at a 90-degree angle. For these measurements, 20 µL DNC dispersion was diluted 100-fold with PBS. Measurements were carried out at the set temperature of 25°C.

Electrophoretic mobility

The superficial charge (ζ-potentials) of the DNC systems was measured using a Zetasizer Nano ZS90 (Malvern). For all samples, an average of three measurements at the stationary level was taken. The temperature was kept constant at 25°C by a temperature controller. The ζ-potential was calculated from the electrophoretic mobility (μζ) using the Henry correction to Smoluchowski’s equation.

Encapsulation efficacy

Free curcumin were removed by means of dialysis. DNC was transferred in a dialysis bag. This dialysis bag was stirred in 360 mL PBS at room temperature for 2 hours. The aqueous medium was refreshed once. The content of curcumin into the dendrosome was quantified by HPLC with diode array detection (HPLC-DAD) analysis, using curcumin as an external standard. The curcumin encapsulation yield is expressed as the so-called encapsulation efficiency (EE), which is defined as the percentage amount of curcumin entrapped in the dendrosome in relation to the total amount of curcumin present during the DNC formation and entrapment procedure.

The curcumin-loaded concentration was assayed by HPLC-DAD analysis performed using the HP 1100 equipped with an HP 1040 DAD, an automatic injector, an auto sampler, and a column oven, and managed by the HP 9000 workstation (Agilent Technologies, Santa Clara, CA, USA). The ultraviolet-visible spectra were recorded between 220–500 nm and the chromatographic profiles were registered at 240, 330, 420, and 430 nm. Separations were performed on a reversed-phase column Luna® RP C18 (150 mm×3 mm id, 5 µm; Phenomenex, Torrance, CA, USA) maintained at 26°C. The eluents were H$_2$O at pH 3.2 by formic acid (A) and acetonitrile (B). The following multistep linear gradient was applied: from 82% A to 56% A in 10 minutes, with a plateau of 3 minutes, in 14 minutes to 48% B, and then 5 minutes to 80% of B. Total time of analysis was 32 minutes, equilibration time was 5 minutes, flow rate was 0.4 mL·min$^{-1}$, and injection volume was 5 µL.

Transmission electron microscopy analysis

A drop (10 µL) of DNC dispersion diluted ten times was applied to a carbon film-covered copper grid (Società Italiana Chimici, Rome, Italy). Most of the dispersion was blotted from the grid with filter paper to form a thin film specimen, which was stained with a phosphotungstic acid solution 1%
weight/volume in sterile water. The samples were dried for 3 minutes and then examined under a JEOL 1010 electron microscope (JEOL, Tokyo, Japan) and photographed at an accelerating voltage of 64 kV.

**Stability studies**

The stability of DNC was studied according to the authors' previous work.26,27 DNC was kept at 4°C±1°C, and at fixed time intervals they were assayed for their physical and chemical stability. Physical stability was checked by monitoring the size and PDI using DLS measurements. Chemical stability was checked by quantification of drug content after disruption of purified vesicles using HPLC-DAD analysis.

**In vitro assessment of the anticancer effect of DNC**

**Cell lines and culture condition**

Glioblastoma cancer cell lines (U87MG), human bone marrow-derived stem cells (hBMSC), and normal human fibroblastic cells (HFSF-P13) were obtained from the national cell bank of the Pasteur Institute, Tehran, Iran. U87MG, hBMSC, and HFSF-P13 cells were cultured in Gibco® RPMI 1640 medium and Gibco® low and high glucose Dulbecco’s Modified Eagle Medium (Life Technologies, Carlsbad, CA, USA), respectively. Cells were supplemented with 10% fetal bovine serum, 100 U/mL penicillin, and 100 mg/mL streptomycin (Life Technologies). All cells were grown at 37°C in a humidified atmosphere of 5% carbon dioxide.

**Cell viability assay**

Cell viability was assessed by methylthiazol tetrazolium (MTT) assay (Sigma-Aldrich) according to the manufacturer’s instruction. Briefly, cells were plated onto 96-well plates, respectively. After 24 hours, the cells were incubated for 24, 48, and 72 hours in the absence or presence of different concentrations of DNC. Dendrosome alone was used to test the cytotoxicity of the nanocarriers. After the treatment, media containing the treatment agents were carefully removed and cells were washed twice with PBS before 20 µL medium containing 5 mg/mL MTT in PBS was added to each well. After 4 hours at 37°C, the medium was totally removed and 200 µL dimethyl sulfoxide solution was added to each well. The absorbance being proportional to cell viability was subsequently measured at 570 nm in each well using an enzyme-linked immunosorbent assay plate reader. All values were compared to the corresponding controls. Cell viability was calculated as the percentage of cell viability of treated cells against control cells.

**Cell cycle analysis**

The effect of DNC on cell cycle distribution was assessed by flow cytometry after staining the cells with propidium iodide (PI). Briefly, 0.3×10⁶ cells (U87MG) were seeded onto six-well plates and allowed to attach overnight. The previous medium was replaced with a fresh one containing the desired concentrations of DNC. After incubating for specified times, cells were collected and washed twice with cold PBS and fixed with ice-cold 70% ethanol at 4°C overnight. After washing twice with cold PBS, cells were resuspended in PBS containing 50 µg/mL PI, 0.1% sodium citrate, and 0.1 Triton X-100 followed by shaking at 37°C for 15 minutes. More than 10⁶ cells were analyzed using a FACSCalibur™ flow cytometer (BD Biosciences, San Jose, CA, USA) and the data were consequently evaluated using Cell Quest software (BD Biosciences). Cells with a lower DNA content than those of the G₀/G₁ phase were referred to as apoptotic cells. The percentage of cells in each phase of the cell cycle was determined by at least three independent tests and the results were expressed as mean ± standard deviation.

**Apoptosis detection**

Annexin-V-FLUOS and PI staining kit (Roche Applied Science, Penzberg, Germany) was used for apoptosis detection following the manufacturer’s instruction. Briefly, U87MG cells were seeded (0.3×10⁶ cells/well) onto six-well plates and allowed to adhere overnight to be treated with desired concentrations of DNC for 12, 24, and 36 hours. Trypsin-digested cells were centrifuged at 200 g for 5 minutes. The cell pellet was resuspended in 100 µL Annexin-V-FLUOS labeling solution and incubated for 10–15 minutes at 15°C–20°C and immediately analyzed by FACSCalibur flow cytometer.

**Caspase activity**

Caspase activity was measured by the homogeneous caspase assay kit (Roche) according to manufacturer’s instructions. Enzymatic activity of caspase-2, 3, 6, 7, 8, 9, and 10 were determined using this assay. Cells were cultured onto 96-well plates (2×10⁴ cell/well) and treated with various concentrations of DNC at different intervals. The amount of free rhodamine 110 resulting from cleavage of the caspase substrate (DEVD-rhodamine 110) was fluorometrically determined at an excitation and emission wavelength of 499 nm and 521 nm, respectively. Data were expressed as relative fluorescence units by Gene 5 software with a fluorescence plate reader (Synergy™ H4 Hybrid Reader;
BioTek, Winooski, VT, USA). Three independent tests were performed and the ratio between the treatment and control group was calculated. Data were expressed as mean ± standard deviation.

RNA extraction and real-time polymerase chain reaction (PCR)
Total RNA was extracted using TRizol® reagent (Life Technologies) followed by DNase I digestion (Thermo Fisher Scientific, Waltham, MA, USA). Complementary DNA was synthesized by PrimeScript™ RT reagent kit (Takara Bio Inc, Shiga, Japan). The list of primers for specific genes and housekeeping gene glyceraldehyde 3-phosphate dehydrogenase (GAPDH) are indicated in Table 1. Real-time PCR was performed using the SYBR® Premix Ex Taq™ II (Takara). Relative gene expression was calculated as 2^-(ΔΔCt).

miRNA expression analysis
The expression of miR-145 and U6 (reference gene) was analyzed using universal specific primer sets and miRCURY LNA™ Universal RT microRNA PCR (Exiqon A/S, Vedbaek, Denmark).

Statistical analysis
Statistics were presented in Prism® 5 software (GraphPad Software, Inc, La Jolla, CA, USA) and analyzed using one-way analysis of variance followed by Newman–Keuls multiple comparison test or Student’s t-test. Differences among groups were stated to be statistically significant when P<0.05.

<table>
<thead>
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<th>Gene</th>
<th>Designed oligonucleotide</th>
<th>Amplicon length (bp)</th>
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<td>OCT4A</td>
<td>F: TCGCAAGCCCTCATTTCC</td>
<td>113</td>
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<td></td>
<td>R: CCATCCCTCCACCATCTC</td>
<td></td>
</tr>
<tr>
<td>OCT4B</td>
<td>F: AGATGGATACTGTTTGAATGTC</td>
<td>191</td>
</tr>
<tr>
<td></td>
<td>R: GCTGAACCATCTCCAAAATGACA</td>
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<tr>
<td>OCT4B1</td>
<td>F: GGTTCTAATTTGGGTTGCC</td>
<td>130</td>
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<tr>
<td></td>
<td>R: TTCTCCCTCTACCTCTC</td>
<td></td>
</tr>
<tr>
<td>SOX-2</td>
<td>F: AAGACTAGACTGAGAGAAGAAG</td>
<td>171</td>
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<tr>
<td></td>
<td>R: AAGAGAGAGCCAACTCAGTGA</td>
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<tr>
<td>Nanog</td>
<td>F: AAATCCCTGCTCAGCAAGAGA</td>
<td>148</td>
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<td></td>
<td>R: TGCCTCAGCACTTGTCTTCT</td>
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<tr>
<td></td>
<td>R: CATGAGTCCCTCCAGATAACC</td>
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</table>

Abbreviations: F, forward primer; R, reverse primer; GAPDH, glyceraldehyde 3-phosphate dehydrogenase; OCT4, octamer binding protein 4; SOX-2, SRY (sex determining region Y)-box 2.

Results
In vitro characterization of DNC formulation
Dimensional analysis, superficial charge, and loading efficiency of DNC
DNC was analyzed in terms of size, PDI, and ζ-potential measured by DLS analysis and in terms of encapsulation efficacy by HPLC-DAD analysis. Regarding DLS measurements, correlation coefficients were superior to 0.9, which means that the signal related to noise ratio was good (10%) (Figure 2A). The mean diameter of DNC was ≤200 nm by distribution algorithm-calculated DLS analysis (Figure 2B and D), and was suitable for systemic administration. PDI is a dimensionless measurement of the broadness size distribution calculated from distribution algorithm, and values were calculated for each peak as peak width/mean diameter. As summarized in Figure 2D, PDI was 0.4; therefore, the colloidal suspension of DNC was polydispersed. The measurements performed three times were nicely reproducible. The ζ-potential was around −7 mV, an acceptable value that does not indicate long-term stability. The curcumin loading efficiency was very high (87%). Transmission electron micrographs by positive stain provided important details on the morphology and dimension of the vesicles. DNC were sphere shaped and had a size comparable with the results from DLS analysis (Figure 2C) considering the different physical principles the two techniques were based on.

Physical and chemical stability tests of DNC
The physical stability of DNC was checked by monitoring size and PDI over time using DLS measurements. Size was maintained within 10 days (Figure 3A), and although PDI increased to 0.6 at day 16, it was constant within 10 days after DNC preparation (0.4). According to these results, no vesicle size alterations or homogeneity loss of the sample occurred within 10 days (Figure 3A). Additionally, chemical stability was checked by quantification of drug content after disruption of purified vesicles. As described in the experimental section, the content of curcumin into the dendrosome was quantified by HPLC-DAD analysis. The chemical stability of curcumin into the dendrosome was maintained at a constant level (not less than 93.8% residual percentage) at least 16 days after their preparation (Figure 3B). Both physical and chemical stabilities (in terms of residual percentage of drug) were acceptable for 10 days.
In vitro assessment of anticancer effect of DNC

DNC induces cell death in U87MG in a time- and dose-dependent manner

The sensitivity of U87MG cells as well as hBMSC and HFSF-PI3 cells to DNC, free curcumin, and empty dendrosome were studied using MTT assay. As shown in Figure 4, DNC significantly suppressed the proliferation of U87MG cells in a time- and dose dependent manner. The half maximal inhibitory concentration (IC<sub>50</sub>) of DNC for U87MG cells was 20 µM after 24 hours (Figure 4A) and 48 hours (Figure 4B), which declined to 10 µM at 72 hours (Figure 4C) (P<0.001). However, the viability of U87MG was not affected by free curcumin only at 72 hours (Figure 4C). On the other hand, no inhibitory effect was observed for dendrosome alone. In total, these findings demonstrate that dendrosome increased the water solubility and entry of curcumin to cells without any toxic effects correlated with dendrosomes as a carrier. Studying the inhibitory effects of DNC on hBMSC showed that these cells are affected by DNC as well as cancerous cells and in a concentration higher than in U87MG cells. Figure 5A shows the effects of DNC on cell viability of hBMSC after 24 hours of drug exposure. Cell viability declined to 67% after treatment with 25 µM DNC (P<0.01) and to 35% and 31% after treatment with 30 µM and 35 µM DNC, respectively (P<0.001). The loss of cell viability was also determined after 48 hours of treatment with DNC (Figure 5B). The cell viability declined to 54% and 52% after treatment with 30 µM and 35 µM DNC, respectively (P<0.05). The viability of HFSF-PI3 was not significantly affected due to treatment and at determined IC<sub>50</sub> for U87MG lines, no inhibitory effect was detected on these cells (P>0.05); however, at 25 µM concentration, the viability of the cells decreased to 50% (Figure 6) (P<0.01). Therefore, in concentrations suppressive for cancer cells, no harmful effects connected to DNC were observed in stem cells and normal fibroblast cells, showing the safety of this formulation as an anticancer treatment agent on normal cells.

The effects of DNC on the U87MG cell cycle distribution

To achieve further insight into the mechanism of the growth inhibitory effects of DNC, U87MG cells were treated in concentr-
concentrations close to the IC$_{50}$ (15, 17.5, and 20 µM) obtained in the MTT assay. Treated cells analyzed for cell cycle distribution were compared to the control. By increasing the concentration of DNC at 24 hours of treatment, the cell population in the SubG$_1$ phase significantly increased ($P=0.0006$). Cells in the G$_1$ or S phase gradually decreased, but not at a statistically significant rate (G$_1$: $P=0.06$; S: $P=0.3$). Surprisingly, cells in the G$_2$/M phase were significantly reduced ($P=0.0147$), consistent with the fact that DNC can target the cell division and proliferation pathways (Figure 7A–C and Table 2). By increasing the time of treatment to 36 hours, an increased cell death rate in the SubG$_1$ phase ($P=0.0036$) together with a gradual decrease in the G$_1$ phase ($P=0.0029$) showed that enhancing the incubation time resulted in probable inhibition of the cell growth pathway (Figure 8). Increasing incubation time from 12 hours to 24 hours led to a significant increase in the SubG$_1$ cell population ($P=0.0215$). However, the rate of cell death slightly decreased at 36 hours compared with 24 hours (Figure 7D, E and Table 3). The cell population in the G$_2$/M and S phases decreased, although these changes were not significant (data not shown). Altogether, these data show that DNC induced cell death in a time- and dose-dependent manner in U87MG cells.

**DNC triggers apoptosis in U87MG cells**

To determine the mode of cell death (apoptosis or necrosis) induced by DNC, Annexin-V-FLUOS and PI staining were applied to separate intact cells, early apoptosis, late apoptosis, and necrotic cells. Different concentrations of DNC (15, 17.5, and 20 µM) were subjected to U87MG after 12, 24, and 36 hours of treatment. As indicated in Figure 9A, DNC induced apoptosis in a time- and dose-dependent manner. Percentages of early apoptosis after 12, 24, and 36 hours of treatment with 17.5 µM were determined as 26.09% ± 0.007%, 30.2% ± 0.02%, and 34.75% ± 0.0%, respectively ($P=0.02$).
Elevated rate of caspase activity in U87MG after DNC treatment

The potential of DNC to activate caspase during apoptosis was determined in a time- and dose-dependent manner. Different concentrations of DNC (15, 17.5, and 20 µM) were subjected to U87MG after 12, 24, and 36 hours of treatment. As illustrated in Figure 9B, DNC induced caspase activation in a time-dependent behavior. Treatment with 17.5 µM DNC resulted in 1.32±0.08-fold, 2.34±0.1-fold, and 3.49±0.009-fold elevated caspase activity in U87MG cells after 12, 24, and 36 hours of exposure, respectively (P<0.0001).

DNC inhibits U87MG cells by silencing the expression of pluripotency transcription factors including OCT4A, OCT4B1, SOX-2, and Nanog

Real-time PCR was used to study the effect of DNC on the expression of pluripotency genes including OCT4A.
variants (A, B, and B1), SOX-2, and Nanog after treatment with 17.5 µM DNC at 12, 24, and 36 hours post-treatment. Designed OCT4 primer sets are potent to discriminate between all OCT4 variants and related pseudogenes. In normal culture conditions, the OCT4A variant is weakly expressed in U87MG and negative expression was observed for the OCT4B variant but OCT4B1, SOX-2, and Nanog were noticeably expressed in U87MG, suggesting that these genes are more important for proliferation in U87MG cells. As shown in Figure 10A, expression of the

![Graph showing cell viability (%) against Concentration (µM) with data points for 24 and 48 hours.](https://www.dovepress.com/)

**Figure 6** Effects of dendrosomal curcumin on the viability of HFSF-P3 cells. Cells were treated with various concentrations of dendrosomal curcumin for 24 and 48 hours. **Notes:** Data expressed as mean ± standard deviation; ***P<0.001 compared to nontreated cells.

![Flow cytometric analysis showing cell cycle distribution in U87MG cells after dendrosomal curcumin treatment.](https://www.dovepress.com/)

**Figure 7** Flow cytometric analysis of the cell cycle in U87MG cells after dose- and time-dependent treatment with dendrosomal curcumin. (A) The cell counts upon treatment with 15, 17.5, and 20 µM dendrosomal curcumin after 24 hours of treatment resulted in the appearance of the SubG1 peak in comparison with control cells. (B) The percentage of cells in the SubG1 phase at different concentrations of dendrosomal curcumin after 24 hours of treatment. (C) The percentage of cells in the G2/M phase at different concentrations of dendrosomal curcumin after 24 hours of treatment. (D) The cell counts upon treatment after 12, 24, and 36 hours of treatment with dendrosomal curcumin. (E) The percentage of cells in the SubG1 phase at different incubation times with dendrosomal curcumin. **Notes:** Data expressed as mean ± standard deviation; *P<0.05, ***P<0.001 compared to control.
Table 2  Cell cycle distribution in U87MG cells after 24-hour dose-dependent treatment of the cells with dendrosomal curcumin using flow cytometry analysis

<table>
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<tr>
<th>DNC (µM)</th>
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<th>Sub-G1 (%)</th>
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<th>G1/M (%)</th>
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</tr>
</tbody>
</table>

Note: Data expressed as mean ± standard deviation.

Abbreviation: DNC, dendrosomal curcumin.

OCT4A variant (P<0.0001), OCT4B1 variant (P=0.0048), SOX-2 (P<0.0001), and Nanog (P=0.007) significantly decreased after 36 hours of treatment with 17.5 µM DNC.

DNC activated miR-145 expression in U87MG cells

The effect of DNC on the expression of miR-145 was investigated. miR-145 is related to the suppression of genes implicated in pluripotency (OCT4 and SOX-2). An increased level of miR-145 can be observed during differentiation. An effect of DNC on miR-145 expression was examined after 24 hours of treatment with 17.5 µM DNC. As shown in Figure 10B, miR-145 expression was elevated approximately 34.35±3.9-fold in comparison with nontreated control cells (P=0.005). The results show that DNC decreased the expression of OCT4A, OCT4B1, and SOX-2 in an miR-145-dependent manner.

Discussion

Glioblastoma is one of the most invasive and heterogenic primary tumors in the central nervous system, with an average survival rate of 1 year. Tumor resection combined with radio- and chemotherapy is the current strategy to treat patients with glioblastoma. The lack of clear margin in glioma tumors results in incomplete resection and tumor recurrence. Additionally, the blood–brain barrier prevents the diffusion of large hydrophobic and ionic molecules to the tumor, making the use of systematically delivered drugs poorly effective. The injection of therapeutic agents into a surgically resected tumor or transient disruption of the blood–brain barrier is a technique suggested to bypass such shortcomings. In spite of the clinical and technical progress in glioblastoma therapy, the mortality rate of this tumor remains high. At present, the increasing knowledge of nanotechnology promises to be helpful in the development of localized delivery of therapeutic agents by means of nanocarriers with small particle size. Here, curcumin was efficiently encapsulated in dendosome micellar structures. The pleiotropic features of curcumin to affect multiple signaling pathways and its preferential targeting of cancer cells rather than normal cells makes it an attractive agent for cancer therapy, and it is safe even at high doses. Given the well-known limitations of glioblastoma treatment strategies, the role of DNC on the proliferation of U87MG—a cellular model of grade IV glioblastoma—was evaluated. The oncogenic characteristics of pluripotency factors including OCT4, SOX-2, and Nanog in glioma cells has been previously reported, feeding into formation of the idea that targeting pluripotency genes may represent a novel approach to providing better treatment for this type of cancer. Here, the expression of two OCT4 variants (A and B), SOX-2, and Nanog, as markers of stemness after treatment with DNC, was measured. Moreover,
Table 3 Cell cycle distribution in U87MG cells after time-dependent treatment of the cells with 17.5 μM dendrosomal curcumin using flow cytometry analysis

<table>
<thead>
<tr>
<th>Cell phase</th>
<th>t=0</th>
<th>t=12 hours</th>
<th>t=24 hours</th>
<th>t=36 hours</th>
</tr>
</thead>
<tbody>
<tr>
<td>SubG₀ (%)</td>
<td>3.9±0.004</td>
<td>37±0.03</td>
<td>56±0.02</td>
<td>47±0.007</td>
</tr>
<tr>
<td>G₁ (%)</td>
<td>39.35±0.005</td>
<td>44±0.1</td>
<td>27±0.006</td>
<td>22±0.004</td>
</tr>
<tr>
<td>S (%)</td>
<td>10.09±0.03</td>
<td>6±0.02</td>
<td>6±0.03</td>
<td>7±0.02</td>
</tr>
<tr>
<td>G₂/M (%)</td>
<td>27.45±0.01</td>
<td>11.75</td>
<td>10±0.03</td>
<td>19±0.01</td>
</tr>
</tbody>
</table>

Note: Data expressed as mean ± standard deviation.
Abbreviation: t, time.

OCT4B1, a novel variant of OCT4 formerly demonstrated by the authors’ research group as a gene involved in the pluripotency pathway, was evaluated. However, the oncogenic roles of OCT4 as a result of related pseudogenes are still controversial. Most of this controversy regarding OCT4 expression is related to OCT4A, which has a strong similarity to OCT4 pseudogenes. The OCT4 primer sets used in the present study are effective in discriminating between all OCT4 variants and related pseudogenes.

In vitro characterization of DNC confirmed that dendrosomes have efficiently encapsulated curcumin in 142 nm spherical structures, which makes it suitable for systemic administration. Analysis of the physical and chemical stability confirmed that DNC has a constant diameter and drug content even 16 days after preparation of this formulation. Data from both MTT assay and flow cytometry show that DNC induced cell death in U87MG cells in a time- and dose-dependent manner, significantly increasing the SubG₀ fraction and reducing the cell population in G₂/M. Together, these data indicate that DNC targets the proliferation pathways in U87MG cells. After 36 hours of exposure the G₁ phase fraction was reduced and the SubG₀ peak was increased, indicating that U87MG growth was inhibited with longer exposure time. Of note, there was no cytotoxicity related to the empty dendrosome in U87MG cells. The effects of DNC in both hBMSC and HFSF-P13 cells were also examined to explore the possible cytotoxic effects on nonneoplastic cells. Calculation of the IC₅₀ values in these cells clearly indicates that cell death occurs in a concentration higher than the one effective in cancer cells. These results may also indicate DNC has preferential uptake in cancer cells relative to normal cells.

Figure 9 (A) Dendrosomal curcumin-induced apoptosis using Annexin-V-FLUOS/propidium iodide (PI) staining in U87MG cells. Viable cells (Annexin-V-/PI-), early apoptotic cells (Annexin-V+/PI-), late apoptotic cells (Annexin-V-/PI+), and necrotic cells (Annexin-V+/PI+) are located in the lower left, lower right, upper right, and upper left quadrants, respectively. Numbers in each quadrant represent the percentage of cells. (B) Caspase-3 activation after treatment with various concentrations of dendrosomal curcumin and different incubation times.

Notes: Data expressed as mean ± standard deviation; **P<0.01; ***P<0.001 compared to control.
as previously mentioned. Protein composition as well as membrane structure diversity between tumor and normal cells could be responsible for drug sensitivity.\textsuperscript{23,32,39} Additionally, DNC can be considered a safe formulation for cancer therapy and clinical applications. Apoptosis depends on caspase activation,\textsuperscript{40} and the role of curcumin in inducing this biologic effect has been investigated in numerous studies that demonstrated its potential to activate different pathways related to apoptosis.\textsuperscript{41,42} In the present study, it was found that DNC-induced cell death in U87MG and was compatible with activation of the apoptotic pathway in a time- and dose-dependent manner. Accordingly, caspase-3 activation was increased by DNC. Previous studies reported that the induction of caspase-3 by curcumin in glioma cells may involve histone hypoacetylation\textsuperscript{43} and activation of the tumor necrosis factor-related apoptosis-inducing ligand (TRAIL/Apo2L) pathway.\textsuperscript{44} Formerly, the role of DNC in apoptosis induction was reported in A431 cells – a cell line model of epidermoid carcinoma – in a PARP (poly [adenosine diphosphate ribose] polymerase-dependent) pathway, indicating that DNC induces cancer cell death via triggering of the apoptosis pathways.\textsuperscript{23} The present study also demonstrates that DNC reduces the expression of stemness genes including OCT4 variants, SOX-2, and Nanog, an action that may limit carcinogenesis. Although Guo et al previously reported that there is a positive correlation between the expression level of OCT4A, SOX-2, and Nanog in glioma and tumor malignancy in a grade-dependent fashion,\textsuperscript{45} this is the first report showing that curcumin has an inhibitory role on OCT4, SOX-2, and Nanog pluripotency genes.

The function of miRNAs as regulators of genes involved in cell growth and proliferation is well established. Among these, mir-145 is one of the best characterized miRNAs in regulation of OCT4, SOX-2, and KLF-4 pluripotency genes. For the first time, Xu et al proved that mir-145 directly targets OCT4 and SOX-2 3'UTR.\textsuperscript{9} Downregulation of mir-145 in a wide range of tumors suggested that this miRNA acts as a tumor suppressor.\textsuperscript{11}

In the present study, miR-145 levels were markedly elevated after treatment with DNC, compatible with the hypothesis that repression of OCT4 and SOX-2 is due to the upregulation of mir-145. Lee et al previously showed that overexpression of mir-145 can significantly inhibit migration and invasion of U87MG/U373MG cells.\textsuperscript{46} Although negative regulation of mir-145 on OCT4 and SOX-2 has been demonstrated in glioma CD133 (+) cells,\textsuperscript{47} the present study shows for the first time that inhibition of OCT4 and SOX-2 by curcumin may occur through mir-145 overexpression. Therefore, miRNA can

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**Figure 10** (A) Effects of dendrosomal curcumin on suppressing pluripotency genes (OCT4A, OCT4B, SOX-2, and Nanog) after treatment with 17.5 µM dendrosomal curcumin after 12, 24, and 36 hours of treatment. (B) Effects of dendrosomal curcumin on the upregulation of miR-145 after 24-hour treatment with 17.5 µM dendrosomal curcumin.

**Notes:** Data expressed as mean ± standard deviation; ***p<0.01; ****p<0.001 compared to control.
be novel targets for natural chemopreventive agents as previously described.48

Conclusion

The novelty of this study is based on the effects of DNC on the pluripotency pathways of glioblastoma cells both at the level of gene expression and regulation. The data show that DNC efficiently suppresses glioblastoma cell growth by downregulation of pluripotency genes, possibly in an miRNA-dependent manner. These results indicate that miRNAs may be used as possible targets for cancer therapy. On the other hand, the lack of DNC cytotoxicity in mesenchymal and fibroblast cells indicates that this new nanodrug can be considered a safe formulation to induce cancer cell death using a natural compound.

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Disclosure

The authors report no conflicts of interest in this work. All the authors have read the manuscript and approved this submission to the *International Journal of Nanomedicine*.

References


Supplementary material

Figure S1 The Fourier transform infrared spectrum of OA400 shows a stretching band of C–H aliphatic at 2,921 cm\(^{-1}\), a stretching band of C–H unsaturated aliphatic at 3,002 cm\(^{-1}\), and a stretching band carbonyl group of ester bands at 1,732 cm\(^{-1}\). The C–H bending vibration of CH\(_2\) and the C–H bending vibration of CH\(_3\) can be seen at 1,462 cm\(^{-1}\) and 1,350 cm\(^{-1}\), respectively. The C–O stretching vibration was at 1,111 cm\(^{-1}\) as a broad band. The C–C stretching vibration can be seen at 1,651 cm\(^{-1}\).