

Anticancer impacts of the unicellular cyanobacterium *Chroococcus turgidus* bioactive compounds in colorectal adenocarcinoma

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Abstract

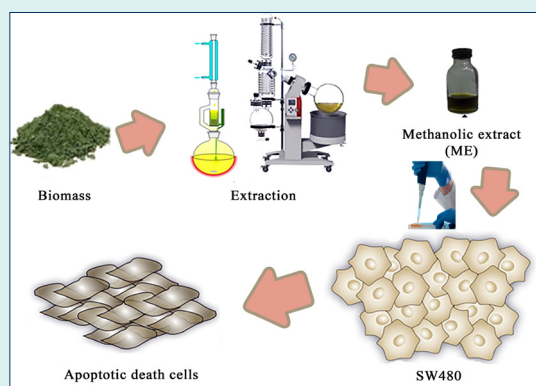
Introduction: Microalgae and cyanobacteria are promising sources of bioactive compounds with antioxidant and anticancer properties. The cyanobacterium *Chroococcus turgidus* has been studied for its potential antioxidant, anti-inflammatory, antibacterial, antiviral, and anticancer effects. This study investigates its anticancer effects on colorectal cancer (CRC) at the cellular and molecular levels.

Methods: The metabolites of *C. turgidus* were screened using the Folin-Ciocalteu reagent and GC-MS.

Antioxidant activity was assessed using the DPPH assay. The biological effects of methanolic extract (ME) were evaluated using MTT assay, Annexin V/PI staining, DAPI staining, and western blotting. Cells were treated with ME at concentrations ranging from 5 to 500 µg/mL for 24 and 48 hours, with the IC₅₀ values determined at 373 µg/mL and 291 µg/mL, respectively.

Results: ME contained bioactive compounds such as phenols, flavonoids, and anthocyanins. Identified fatty acids included palmitic acid ethyl ester (15.53%), 1-bromo-11-iodoundecane (2.31%), undecanoic acid 2,8-dimethyl methyl ester (6.62%), oleic acid (6.47%), and 7-dehydrocholesterol (7.97%). ME inhibited SW480 cell proliferation in a dose- and time-dependent manner and induced nuclear fragmentation, chromatin remodeling, and apoptosis. Annexin V/PI staining confirmed apoptosis as the dominant mode of cell death. Western blot analysis showed increased Bax and decreased Bcl2 expression, supporting its pro-apoptotic activity.

Conclusion: *C. turgidus* may serve as a potential therapeutic agent for gastrointestinal cancers through its ability to modulate the Bax/Bcl2 pathway and promote apoptosis. These findings highlight its novel anticancer effects and support further preclinical investigations.



Introduction

In recent years, cancer has emerged as one of the most significant global health challenges, consistently ranking as the second leading cause of death across many nations. The World Health Organization (WHO) anticipates

a rise in cancer-related mortality rates, particularly in low- and middle-income nations, where access to early detection, treatment, and healthcare resources remains limited. This growing burden underscores the urgent need for comprehensive prevention strategies, improved



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healthcare infrastructure, and equitable access to cancer care worldwide.¹ Globally, colorectal cancer (CRC) ranks as the second most common cancer among women and the third among men.² The mortality rate of CRC may vary by as much as tenfold, highlighting the widening disparities and increasing disease burden in countries undergoing economic and healthcare transitions. While significant progress has been made in improving CRC outcomes in more developed nations, there persists a significant clinical gap, particularly in regions with limited resources, where access to early detection, advanced treatments, and comprehensive care is often restricted. Addressing these gaps is crucial for reducing the global burden of CRC and ensuring equitable healthcare outcomes.^{3,4} Different approaches are routinely used in the treatment of various cancers, including chemotherapy, radiotherapy, surgery, cryosurgery, radiation therapy, and immunotherapy.⁵ However, these treatment modalities are often associated with expensive processes and may result in severe side effects.^{6,7} Moreover, these combined treatment approaches are not without flaws, often leading to various secondary health threats, unspecific outcomes, and unavoidable toxicity.^{8,9} In light of these challenges, there is an urgent demand for cost-effective treatment strategies that exhibit reduced side effects and are suitable for a broad range of cancers. The use of traditional natural product-based strategies is becoming more widely recognized for their potential in pharmaceutical innovation, offering promising solutions. The successful isolation and development of natural biomaterials (e.g., carbohydrates, lipids, proteins, enzymes, and secondary metabolites) have opened new avenues for the creation of therapeutic compounds and pharmacophores. This progress has invigorated the scientific community, driving renewed efforts to harness these natural products in the drug discovery and evaluation process with greater enthusiasm and focus.^{8,10-12} Significant efforts have been dedicated to isolating such compounds, leading to the identification of over 10,000 natural products that may have biotechnological applications. However, the overwhelming abundance of metabolites and their vast dynamic range have limited access to many bioactive natural products, hindering further investigation. Additionally, complex challenges (e.g., evaluating pharmacokinetics, pharmacodynamics, and safety parameters) have emerged as major concerns in the study of these natural products.¹³⁻¹⁵ The unique ecological, chemical, and biological properties of marine environments have endowed microalgae and cyanobacteria with the potential to produce a variety of bioactive chemical compounds. These compounds, such as phenols, flavonoids, and anthocyanins, have various health benefits and find applications in therapeutics.¹⁶ Notably, cyanobacteria, traditionally known for containing chlorophyll a and phycobiliproteins (i.e., phycocyanin and phycoerythrin), have been found to exhibit greater

pigment diversity. Recent research suggests the presence of chlorophyll b in some cyanobacteria, challenging previous notions and indicating an evolutionary complexity in their photosynthetic apparatus.¹⁷ This includes potential ancestral traits and adaptations allowing efficient light capture across different environments. Such findings reveal that cyanobacteria can utilize various chlorophylls (including chlorophylls d and f), enhancing our understanding of their photosynthetic mechanisms and evolutionary history.^{18,19} Microalgae and cyanobacteria-derived phytochemicals have distinct and high potential biological actions compared to the phytochemical constituents of terrestrial origin (plant phytochemicals).^{16,20,21} Of cyanobacteria, some species have been approved as safe for human consumption, such as *Spirulina* or *Chlorella*, in large part due to their useful secondary metabolites. Notably, lower organisms elaborate numerous secondary metabolites or natural products as signaling molecules for "offense and defense". Recently, these metabolites have been extracted, their actions were checked in various bioassays, and their potential as a remedy for human diseases was evaluated.²²⁻²⁴ This bioprospecting study aimed to develop novel anticancer agents with enhanced efficacy, focusing on the unicellular cyanobacterium *Chroococcus turgidus*, which was isolated from the KANI Barazan International Wetland, located to the south of Lake Urmia. The methanolic extract (ME) of *C. turgidus* was evaluated for its potential to induce apoptosis in the SW480 colon cancer cell line. The antioxidant activity of the ME was assessed using the diphenyl picryl hydrazyl (DPPH) assay. In addition, its ability to inhibit cancer cell growth was thoroughly investigated through multiple techniques, including flow cytometry, DAPI (4',6-diamidino-2-phenylindole) staining, and western blot analysis. The combination of these methods facilitated a comprehensive evaluation of the ME's anticancer properties.

Materials and Methods

Materials

Ascorbic acid, Folin-Ciocalteu reagent, and Quercetin were purchased from Sigma-Aldrich (St. Louis, MO, USA). Gallic acid, Methanol, and DPPH were obtained from Merck (Kenilworth, NJ, USA). The human colorectal carcinoma SW-480 cell line was obtained from the National Cell Bank of Iran, Pasteur Institute (Tehran, Iran). RPMI 1640 medium, fetal bovine serum (FBS), and trypsin-EDTA (0.02–0.05%) were acquired from Gibco (Paisley, UK). Phosphate-buffered saline (PBS) was purchased from Sigma-Aldrich Company (Munich, Germany). Mouse β -Actin (sc-47778) and Bax (sc-7480) monoclonal antibodies, rabbit Bcl2 (sc-492) antibody, and mouse anti-rabbit IgG-HRP (sc-2357) were obtained from Santa Cruz Biotechnology (Santa Cruz, CA, USA). The reverse transcriptase reagent and FITC-labeled

annexin V- apoptosis detection kits were purchased from TAKARA Co. (Tokyo, Japan) and Applied Biosystems (Foster City, CA, USA), respectively.

Isolation of *C. turgidus*

The water samples were collected from the regional KANI Barazan International Wetland, which located 30 km from Mahabad city in the West Azerbaijan province (NW Iran) with the geographical position of N 36° 59' 34" and E 45° 46' 34". The identification of *C. turgidus* was accomplished by using the algal flora keys. For the purification of *C. turgidus*, single colonies were picked and transferred into the 50 mL flasks containing 1M modified BG 11's medium composed of NaNO₃ (. 15 g/L), K₂HPO₄ (4.0 g/L), MgSO₄.7H₂O (7.5 g/L), CaCl₂.2H₂O (3.6 g/L), C₆H₈O₇ (0.6 g/L), (NH₄)₅[Fe(C₆H₄O₇)₂] (0. 6 g/L), EDTANa₂ (0.1 g/L), Na₂CO₃ (2.0 g/L), H₃BO₃ (2.86 g/L), MnCl₂.4H₂O (. 181 g/L), ZnSO₄.7H₂O (0.22 g/L), Na₂MoO₄.2H₂O (0.39 g/L), CuSO₄.5H₂O (0.08 g/L), Co (NO₃)₂.6H₂O (0.05 g/L). Moreover, the culture media of *Chroococcus turgidus* was kept under the required condition (i.e., 26 °C temperature with 16:8 light: dark photoperiod, and 80 μmol photon m⁻² s⁻¹ irradiance).

Molecular characterization of *C. turgidus*

After morphological identification, molecular characterizations using 16S rRNA gene and 16S-23S ITS region were applied for the accurate and reliable identification of the isolated cyanobacteria, identified as *C. turgidus* strain KANI.²⁵

Preparation of methanolic extracts of *C. turgidus*

The cyanobacteria cells were harvested from 500 mL of BG 11's medium (O.D. 630_{nm}. 15–2.0) by centrifugation (3000g for 5 minutes, at 4 °C) and then freeze-dried for the next process. The soxhlet apparatus extracted freeze-dried cyanobacteria biomass (5 g) with 125 mL of solvent for 7 hours. The extracts were concentrated in a Buchi rotary evaporator at 120 rpm and 60 °C for 2 hours, and then traces of solvent were removed using a desiccator. After filtration of 3 mL ME through the Whatman paper (grade 42), the gas chromatography-mass spectrometry (GC-MS) analysis was performed. The samples were dried under a laminar flow hood, and absolute alcohol and sodium sulfate were added to remove residual water. A quality control sample (blank solvent) was included to ensure the absence of contamination. The GC-MS analysis was conducted using an Agilent 6890 equipped with an HP-5MS (5% diphenyl/95% dimethyl polysiloxane) fused silica capillary column (30 m × 0.25 mm i.d., film thickness 0.25 μm). The compounds were identified by comparing the obtained mass spectra with those available in the National Institute of Standards and Technology (NIST) library, which contains over 62,000 reference patterns. For further validation, an internal standard (n-alkane series,

C10 – C40) was used to calibrate retention times, and the spectra of unknown compounds were matched with authenticated reference spectra from the NIST database. Additionally, retention indices (RI) were calculated and compared with literature data to enhance the reliability of compound identification.

Analysis of photosynthetic pigments

The photosynthetic pigments, including chlorophyll *a*, *b*, and total carotenoids, were assessed based on the previously described method.²⁶ Briefly, cyanobacteria cells were harvested from 5 mL of media (O.D. 630_{nm}. 15–2.0) by centrifugation (at 26 °C, 3000 g for 5 minutes). The collected biomass (equivalent to approximately 15 mg dry weight) was homogenized in 5 mL of 100% methanol at 4 °C for 24 hours in the dark. Then, the homogenates were centrifuged at 10,000 × g for 10 minutes at 4 °C to remove cell debris. For quantitative determination of pigments, the supernatants were analyzed using a UV/V spectrophotometer at 470, 665, and 653 nm. Methanol was used as a blank to correct baseline absorbance, and potential spectral interferences were minimized by ensuring complete extraction of pigments and avoiding contamination from cellular debris. The content of the pigments was determined based on standard equations. The content of the pigments was determined,^{26,27} as follows:

$$C_a = 15.65 A_{665} - 7.340 A_{653}$$

$$C_b = 27.05 A_{653} - 1.121 A_{665}$$

$$C_{x+c} = (1000 A_{470} - 2.860 C_a - 129.2 C_b) / 245$$

Where, *A* denotes the absorbance in the presence of the sample, *C_a* and *C_b* represent the chlorophyll *a*, *b* and *C_{x+c}* denotes the total carotenoids in the presence of the sample.

Analysis of total phenol, flavonoid contents

Total phenols and flavonoids of the cyanobacteria cells were isolated using 100% methanol. The total phenol quantity of the extracts was measured using the Folin-Ciocalteu procedure according to the previous method.^{28,29} In brief, 100 μL of cyanobacterial extract was mixed with 2.8 mL of deionized water, 100 μL of Folin-Ciocalteu reagent, and 2 mL of sodium carbonate aqueous solution (2%, final concentration 0.4%). The sodium carbonate solution was prepared by first dissolving 2 g of NaOH in 500 mL of distilled water to obtain a 0.1 M NaOH solution. Then, 2 g of Na₂CO₃ was dissolved in 100 mL of this NaOH solution. The reaction mixture was incubated in the dark at room temperature for 30 minutes. Absorbance was measured at 720 nm using a UV/Vis spectrophotometer, with methanol as the blank. A calibration curve was prepared using gallic acid as the

standard, and results were expressed as mg gallic acid equivalent (GAE) per gram of fresh weight (mg GAE/g F.W.).

The total flavonoid content was measured using an aluminum chloride colorimetric assay. For each reaction, 500 µL of cyanobacterial extract was mixed with 15 mL of 100% methanol, 100 µL of freshly prepared aluminum chloride solution (10%) (prepared by dissolving 10 g of AlCl₃ in 90 mL methanol), 100 µL of potassium acetate (1 M), and 2.8 mL of distilled water. The mixture was incubated at 25 °C for 40 minutes. The absorbance was recorded at 415 nm using a spectrophotometer, with methanol as the blank. The flavonoid content was calculated as mg quercetin equivalent per gram of fresh weight (mg QE/g F.W.).

For anthocyanin quantification, 500 µL of the extract was mixed with 4.5 mL of acidified methanol (1% HCl in methanol, v/v), and the mixture was incubated at 4 °C for 24 hours in the dark to prevent pigment degradation and enhance extraction efficiency. After incubation, the sample was centrifuged at 10,000×g for 10 minutes at 4 °C. Absorbance was measured at 530 nm and 657 nm, with methanol as the blank, and the total anthocyanin content was calculated using the equation:

$$\text{Total Anthocyanin (mg/L)} = (A_{657} \times 0.25 - A_{530})$$

Where A_{657} and A_{530} correspond to the absorbance at 530 and 657 nm, respectively. Cyanidin-3-glucoside was used as the standard for quantification.

Analysis of anthocyanin content

A 20 mg cyanobacteria cells sample was crushed in a porcelain mortar with 4 mL of hydrochloric acid containing 1% methanol to determine the total anthocyanin content. The mixture was kept in the refrigerator for 24 hours and then centrifuged at 13000×g for 10 minutes. After collecting the supernatant, the absorbance (A) of the solution was determined at 530 and 657 nm and normalized against the control sample of hydrochloric acid containing 1 % methanol. The absorbance used for quantifying the anthocyanin content was calculated using the following equation.³⁰

$$A = A_{530} - (0.25 \times A_{657})$$

Where A denotes the absorbance in the presence of the sample.

Free radical DPPH scavenging capacity

The DPPH radical scavenging assay was carried out based on a method described previously by Ozturk and Tuncel.³¹ The DPPH assay is a free radical method based on the radical scavenging activity of antioxidants towards the purple-colored DPPH in methanol. The hydrogen donors of antioxidants can reduce the free radical DPPH to the corresponding stable diamagnetic molecule hydrazine

(yellow-colored).^{31,32} Reaction mixtures of samples were prepared by mixing appropriate amounts of extract with different concentrations (100, 200, 300, 400, and 500 µg/mL), 2 ml of DPPH (0.1 mM in methanol 96%) to a total volume of 4 mL. All samples vortexed (1 min) and incubated in the dark for 60 min at 37 °C. All the experiments were done in triplicate. DPPH (0.1mM) was taken as control and ascorbic acid as standard. The decrease in absorbance of each sample was measured against methanol as blank on a spectrophotometer, Ultraspec 2000 (Pharmacia Biotech Co., Garden City, England) at 517 nm. The percentage of DPPH was calculated using the following equation:

$$\text{DPPH\%} = \frac{A_c - A_s}{A_c} \times 100$$

Where, A_c and A_s denote the absorbance of the control reaction and the absorbance in the presence of the sample, respectively.

Cytotoxicity assays

The SW-480 cells were cultivated at a seeding density of 10×10^4 cells/well in 96-well plates. After 24 h, the cells were exposed to 200 µL of fresh medium containing various concentrations (5, 10, 50, 100, 200, 300, 400, and 500 µg/mL) of the extracts. These concentrations were selected based on preliminary dose-response studies. After 24 and 48 hours, the medium of each well was replaced with 200 µL of fresh MTT [3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide] reagent (2 mg/mL) for 4 hours. Next, the MTT reagent was removed, and then 200 µL dimethyl sulfoxide (DMSO) was added to stop the reaction, and the cells were incubated at 37 °C for an additional 15 min. The optical density was determined using a microplate reader, ELx808 (BioTeck, Winooski, VT), at 570 nm wavelength. The viability of cells was evaluated relative to the absorbance of untreated control cells. All experiments were carried out in triplicates.

Apoptosis assay by Annexin-V

A flow cytometry assay was performed to determine the induction of apoptosis in the treated SW480 cells with microalgae extract. Briefly, the SW480 cells were treated with ME (IC₅₀ Dose) and incubated at 37°C for 24 hours. The final concentration of cells before staining was (10×10^5 cells/well in 12-well plates). The cells were then resuspended in 200 µL of annexin V-binding buffer (from EXBIO) and incubated at room temperature in the dark for 10 min. Afterward, the cells were washed with ice-cold PBS and subjected to flow cytometry analysis using FACSCalibur™ (Becton Dickinson Co., Franklin Lakes, NJ, USA) with an emission filter of 600 nm for P.I. and 515–545 nm for FITC.³³ To ensure the validity of the analysis, untreated cells were used as a control group.

Apoptosis assay by DAPI staining

Given that nuclear fragmentation and chromatin condensation and remodeling are the typical markers

of apoptosis, the DAPI staining assay was employed to analyze the occurrence of such phenomena in the treated cells. Briefly, the SW480 cells were treated with ME (IC₅₀ Dose) and incubated at 37°C for 24 hours. Next, the cells were fixed with the freshly prepared ice-cold paraformaldehyde 4%. Then, to permeabilize the cells, they were exposed to 0.1% Triton X-100 in PBS (NaCl, KCl, KH₂PO₄, and Na₂HPO₄, pH 7.4) for 5 minutes. The cells were stained with DAPI (1 µg/mL in PBS) in the dark for 3 min. Afterward, the cells were washed (×3) with 0.1% Triton X-100 in PBS. Next, they were assessed using a live imaging system, Cytation™ 5 (BioTek, Winooski, USA).

Western blotting assay

The SW480 cells were cultivated at a seeding density of 2.0×10^5 cells/well in 6-well plates. Briefly, the SW480 cells were treated with the IC₅₀ dose of ME and incubated at 37 °C for 24 hours. The treated and untreated cells were centrifuged to extract proteins using radioimmunoprecipitation assay (RIPA) lysis buffer containing a protease inhibitor cocktail (PMSF, leupeptin, and aprotinin). The homogenate was centrifuged at $5000 \times g$, 4 °C for 5 minutes. The concentration of protein samples was determined using the Bradford assay with BSA as the standard. Equal amounts of total protein were loaded per well of a 12% polyacrylamide gel (SDS-PAGE). The proteins were then transferred onto a polyvinylidene fluoride (PVDF) membrane at 100 V for 1 hour. The membranes were incubated in 5% bovine serum albumin (BSA) dissolved in 20 mM Tris-HCl, containing 150 mM NaCl and 0.05% Tween-20 at 4 °C overnight. After three washes, the membranes were incubated with specific primary antibodies (1:4000; Bax, Bcl2, and β-actin) followed by the secondary antibody (1:8,000; horseradish peroxidase-conjugated) in 3% BSA at room temperature for 2 hours and 1 hour, respectively. Protein bands were detected using the Pierce™ ECL western blotting substrate chemiluminescent kit. The expression level of β-actin was used to normalize the protein levels. The quantification of Bax and Bcl2 expression was compared with the expression of the housekeeping protein and analyzed using ImageJ software.

Statistical analyses

The data were analyzed to determine statistical differences using one-way ANOVA followed by Tukey's post hoc test. Statistical analyses were performed with SPSS software version 19.0, and a *P* value of ≤ 0.05 was considered statistically significant. All data are presented as mean \pm standard deviation (SD).

Results

Morphological and molecular traits of isolated cyanobacteria

To identify the collected cyanobacteria, a valid authentication key, algal flora, and some recent related

references were carefully used.^{34,35} *C. turgidus* strain KANI was identified based on some morphological characteristics, including shape and isolated two- or four-celled groups together with an amorphous mucilage sheath (Fig. 1). For the confirmation of morphologic findings, the 16S rRNA gene and 16S-23S ITS region from the cyanobacteria genome were amplified. Consequently, the samples were sequenced and blasted by means of the National Center for Biotechnology Information (NCBI) database. The blasting data confirmed the morphologic results where the isolated cyanobacteria showed a high genetic similarity to *C. turgidus*.

Fatty acid identification by GC-MS analysis

GC-MS analysis indicated the occurrence of different compounds, especially fatty acids, including fatty acid esters (FAEs), oleic acid, palmitic acids, and provitamin D3 in *C. turgidus* strain KANI ME (Table 1).

Spectrophotometric analysis of photosynthetic pigments microalgae

Chl *a*, Chl *b* and carotenoid levels were determined with a UV-Vis spectrophotometer measuring absorbencies at 665, 649 and 470 nm, respectively, as shown in Fig. 2a. Besides, the level of photosynthetic pigments in the extract was measured for chl *a* and *b*, and carotenoids were 18.424, 12.864, and 9.428 µg/g FW, respectively (Fig. 2b).

Estimation of phenol, flavonoid, and anthocyanin contents

As shown in Fig. 3, the ME 's phenols, flavonoids, and anthocyanins content were measured at a concentration of 7.428, 1.1864, and 4.424 mg/g FW, respectively. Based on the results, flavonoids are the dominating phenolic compounds in *C. turgidus* strain KANI.

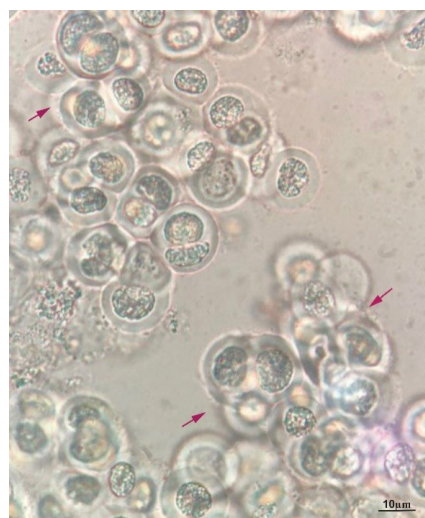


Fig. 1. The morphology of the isolated *C. turgidus* strain KANI. Red arrows depict two, four, and eight-celled colonies enclosed with an amorphous (Scale bar = 10 µm).

Table 1. The identified fatty acids in *Chroococcus turgidus* strain KANI by GC-MS

Compound analyzed	Retention Time	Molecular Formula	Molecular weight	Peak area%	Nature of compound	Bioactivity
1-Bromo-11-iodoundecane	24.54	C ₁₁ H ₂₂ BrI	36.11	2.31	Halogenated fatty acid	Antimicrobial, Antifungal ³⁶
Palmitic acid, ethyl ester	24.70	C ₁₈ H ₃₆ O ₂	284.5	15.53	Fatty acid ester	Antioxidant, Nematicide, Insecticide, Lubricant, Hemolytic, Hypocholesterolemic, Pesticide, Antiandrogenic, Flavor, Hemolytic, 5-Alpha reductase inhibitor ³⁷⁻³⁹
Undecanoic acid, 2,8-dimethyl-, methyl ester	24.86	C ₁₄ H ₂₈ O ₂	228.37	6.62	Fatty acid ester	Antibacterial, Antitumor ⁴⁰
Oleic acid	27.04	C ₁₈ H ₃₄ O ₂	282.5	6.47	Fatty acid	Antitumor, Lubricant Antibacterial, Antitumor, antioxidant and Anticancer Antimicrobial, Antiandrogenic ^{41, 42}
7-Dehydrocholesterol	29.46	C ₂₇ H ₄₄ O	384.6	7.97	Fatty acid	Provitamin D3 ⁴³

Determination of antioxidant activity

For the analysis of the antioxidant activity, the EC₅₀ is determined from the dependence between the DPPH concentration remaining after its reaction with the antioxidant and different antioxidant concentrations. As a result, the EC₅₀ value was estimated at around 400 µg/mL of the extract (Fig. 4).

Evaluation of the cytotoxic effect of ME

The ME concentrations from *C. turgidus* strain KANI were evaluated for their cytotoxic effects on SW480 cancer cells using the MTT assay. As shown in Fig. 5, viable cell numbers were significantly reduced ($P < 0.05$) in a dose- and time-dependent manner, with the greatest effect observed at 48 hours. The statistical analysis confirmed that both dose and time had a significant impact on cell viability. The IC₅₀ values for ME after 24 and 48 hours were 373 µg/mL and 291 µg/mL, respectively. Thus, it can be deduced that the cyanobacterial extract affects SW480 cells through a dose- and time-dependent mechanism (Fig. 5).

Evaluation of apoptosis by Annexin V

To analyze the apoptosis in SW480 cells treated with ME (IC₅₀ Dose) using Annexin-V, PI was removed due to its spectrum overlap (620 nm) with photosynthetic pigments spectrum in the ME (600-700 nm). In comparison

with the untreated control cells, after 24 hours, 38.97% apoptosis was induced in the SW480 cells treated with the IC₅₀ concentration of the ME (Fig. 6).

Apoptosis evaluation by DAPI staining

DAPI is a cell-permeable fluorescent compound that stains DNA by binding with high affinity to the minor groove at A-T-rich regions. As shown in Fig. 7, compared to untreated cells, a notable proportion of treated SW480 cells exhibited apoptotic nuclear features, including chromatin condensation and nuclear fragmentation, following 24-hour exposure to ME at the IC₅₀ dose. These morphological changes are characteristic of apoptosis and further support the cytotoxic effects of the extract.

Apoptosis induction by protein expression assay in Western-blotting

As shown in Fig. 8, treatment with ME (IC₅₀ Dose) has reduced the expression of *Bcl2* proteins and increased the *Bax* expression in the SW480 cell line. In addition, the *Bax/Bcl2* ratio was significantly different between the treated and control groups, indicating the relative protein expression of *Bcl2* and *Bax* compared to the control group.

Discussion

Over the past few decades, more than 50 000 natural products have been discovered from marine

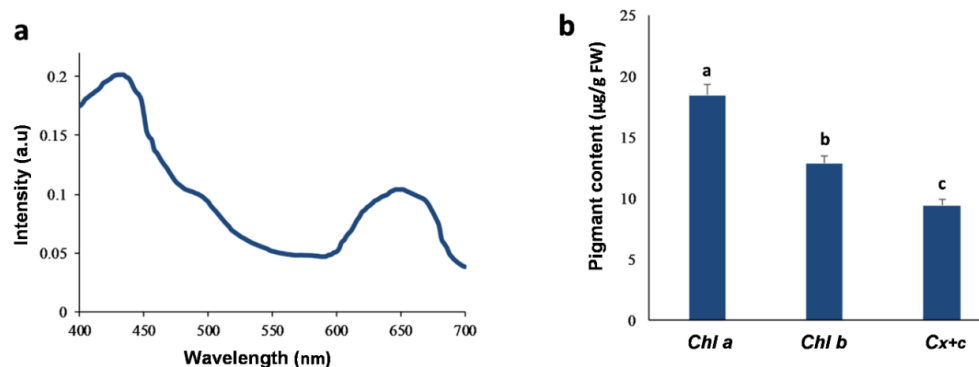


Fig. 2. Photosynthetic pigments from the methanolic extract (ME) of *C. turgidus* strain KANI. (a) visible absorption spectra of ME (b) Chlorophylls (Chl a and b) and total Carotenoids (Cx+c) contents. Columns labeled with different letters are significantly different ($P < 0.05$). Data were expressed as mean values of independent triplicates (mean \pm SD).

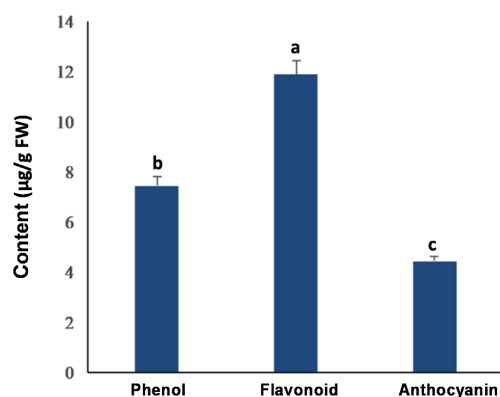


Fig. 3. Total phenol, flavonoid, and anthocyanin contents from the methanolic extract (ME) of *C. turgidus* strain KANI. Columns labeled with different letters are significantly different ($P < 0.05$). Data were expressed as mean values of independent triplicates (mean \pm SD).

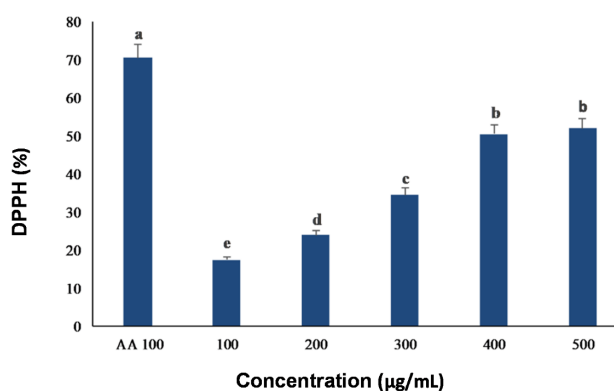


Fig. 4. DPPH radical scavenging activity from the methanolic extract (ME) of *C. turgidus* strain KANI compared with standard (AA; ascorbic acid). Data were expressed as mean values of independent triplicates (mean \pm SD). Different subscripts in small letters (a, b, c, d, and e) above the columns indicate significant differences at $P \leq 0.05$.

microorganisms, many of them with biomedical applications.⁴⁴⁻⁴⁶ Research on molecules produced by aquatic organisms reveals that microalgae and cyanobacteria synthesize a vast array of compounds with promising biotechnological applications. Notably, these compounds exhibit significant therapeutic potential against cancer cells due to their diverse biological functions. These include antioxidant activity, anti-inflammatory properties, anti-mutagenic effects, inhibition of cell proliferation, promotion of cell cycle arrest, induction of apoptosis or autophagy, potential anti-invasion and anti-metastasis effects, suppression of drug resistance mechanisms, and enhancement of chemotherapy sensitivity.⁴⁷⁻⁵⁴ The growing interest in marine-derived natural products has led to the identification of bioactive flavonoids, polyphenols, and sterols with potent anticancer activity. Notably, compounds such as quercetin and resveratrol, widely recognized for their anti-tumor properties, exert their effects through oxidative stress modulation, PI3K/AKT inhibition, and apoptotic signaling activation.^{55,56}

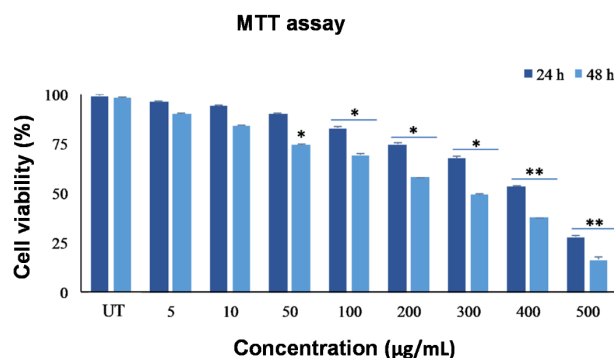


Fig. 5. Cell viability assay of the SW480 cell line in different concentrations (0-500 µg/mL) from the ME of *C. turgidus* strain KANI. Data were expressed as mean values of independent triplicates (mean \pm SD). Indicates significant difference in compared to control group (* $P < 0.05$ and ** $P < 0.01$). ME: methanolic extract, UT: untreated control.

To contextualize our findings, we have expanded our discussion by comparing the anticancer effects of *C. turgidus* metabolites with these well-characterized plant-derived compounds, highlighting the unique mechanistic aspects of our study.

Fucosterol, a sterol commonly found in marine algae, has demonstrated anti-inflammatory, antioxidant, and anticancer properties, particularly against hematologic malignancies.⁵⁷ Moreover, hexadecanoic acid methyl esters were highlighted for use as anti-inflammatory agents, cancer preventive, hepatoprotective, antiarthritic, and anti-coronary attributes.⁵⁸⁻⁶⁰ In our study, we identified 7-dehydrocholesterol (provitamin D3) as one of the major components of *C. turgidus* extract. Given the established link between vitamin D deficiency and increased CRC risk, this sterol suggests an additional potential mechanism of action, warranting further investigation into its role in modulating cancer cell behavior.⁶¹ To explore new anticancer molecules with potentially fewer side effects and reduced resistance to existing drugs, this study focused on isolating and identifying compounds from cyanobacteria in the Kani Barazan International Wetland, located to the south of Lake Urmia. The cyanobacterium *C. turgidus* strain KANI was successfully isolated from the wetland, with its microscopic morphology aligning with the Chroococcaceae family. Molecular identification was performed using the 16S rRNA gene and the 16S-23S ITS region from its genome. Based on NCBI database recommendations, the organism was given a Barcode of Life identifier (MW040530.1, 2021)_ENREF_49.²⁵ The amount of natural compounds in *C. turgidus* strain KANI is very impressive, which increases the importance of this cyanobacterium for the cultivation and extraction of these compounds for medicinal purposes. Other related investigations have reported the therapeutic effects of the identified fatty acids in this study (Table 1).³⁶⁻⁴³

The cytotoxicity assay results demonstrated that the methanolic ME of *C. turgidus* strain KANI exerted a dose- and time-dependent inhibitory effect on the proliferation

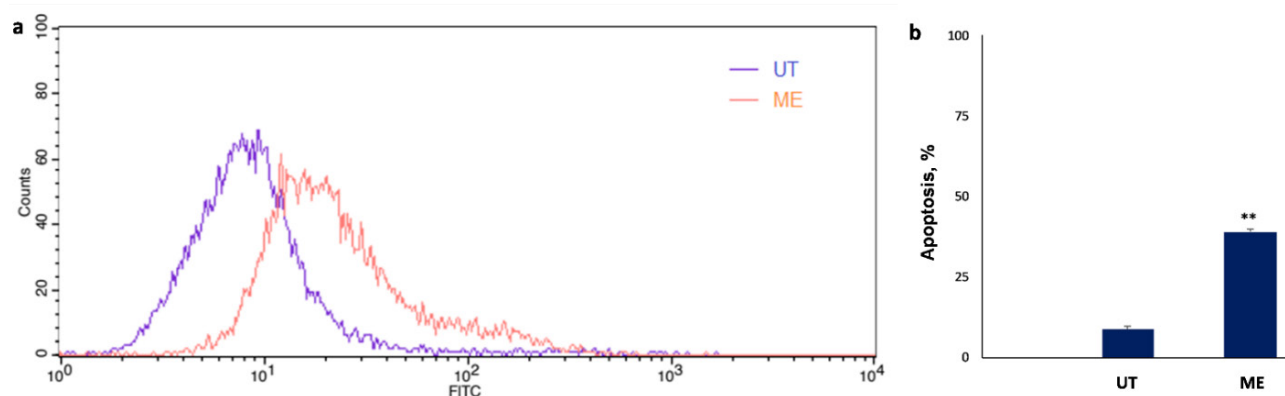


Fig. 6. Apoptosis assay using Annexin-V flow cytometry analysis on SW480 cancer cells. (a) Treated cells with ME comparison with untreated control cells. (b) The rate of apoptosis in the treated cells with ME data was expressed as mean values of independent triplicates (mean \pm SD). Indicates a significant difference compared to the control group (** $P < 0.01$). ME: methanolic extract, UT: untreated control cells.

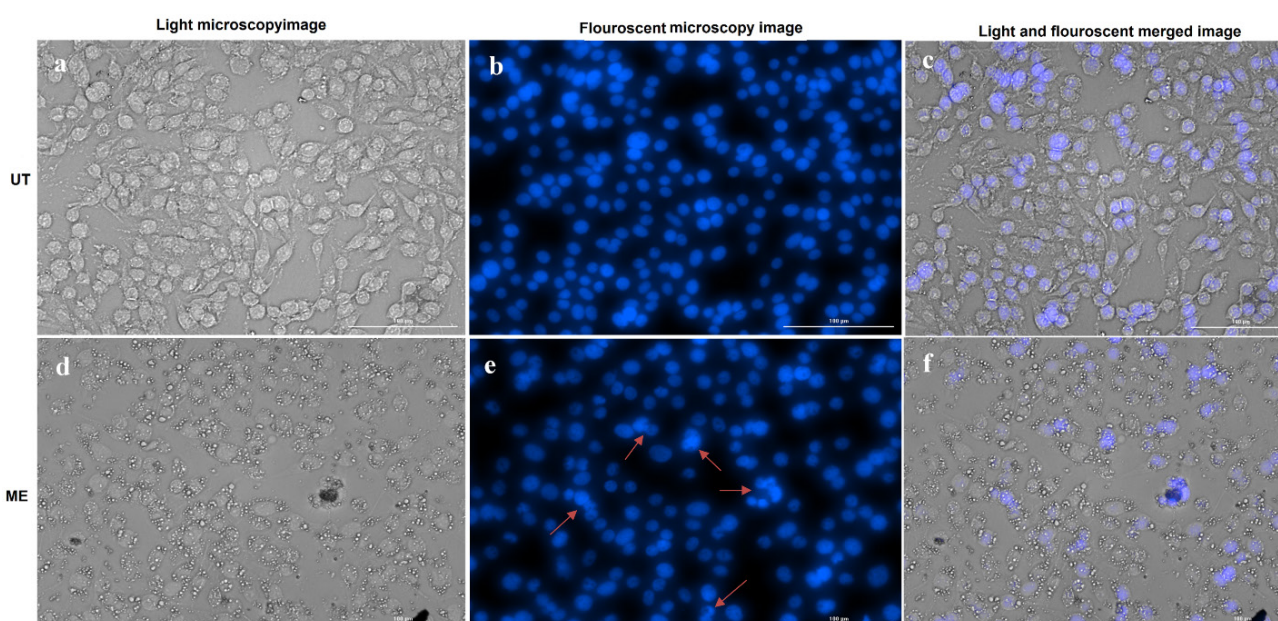


Fig. 7. Apoptosis assay by DAPI staining of SW480 cells. (a, b and c) UT group and (d, e, and f) treated with the ME. Red arrows depict chromatin condensation and fragmented nuclei. ME: methanolic extract, UT: untreated control cells.

of SW480 CRC cells, consistent with previous reports on the pro-apoptotic effects of microalgal metabolites. The cytotoxic effects suggest that *C. turgidus* metabolites primarily exert their anticancer activity through apoptosis induction.^{62,63} However, the therapeutic potential of these compounds extends beyond their in vitro cytotoxicity and is influenced by their bioavailability and metabolic stability. Several studies have highlighted that the pharmacokinetics of marine-derived bioactive compounds can significantly impact their therapeutic applicability.^{64,65} While this study focused on in vitro cytotoxic effects, further *in vivo* validation is necessary to evaluate these metabolites' absorption, distribution, metabolism, and excretion properties. Structurally similar compounds, such as polyphenols and sterols, exhibit variable bioavailability due to metabolic modifications; therefore, future research should focus on optimizing

formulation strategies that enhance the stability and systemic delivery of compounds originated by *C. turgidus*, potentially through nanoscale delivery systems.

Mechanistically, natural compounds can exert cytotoxic effects through multiple pathways. For instance, curcumin and curcuminol induce apoptosis via p53 activation, oxidative stress modulation, and NF- κ B suppression, while kaempferol inhibits angiogenesis through the ERK/NF- κ B/c-Myc axis. Moreover, zerumbone, a naturally occurring sesquiterpene compound found primarily in the rhizomes of *Zingiber zerumbet*, promotes apoptosis by enhancing the expression of pro-apoptotic proteins such as Bax via cytochrome c release and activating caspase cascades, while simultaneously reducing the levels of anti-apoptotic proteins like Bcl2.⁶⁶⁻⁶⁹ There is increasing evidence to prove the prognostic and predictive role of apoptosis-related markers such as Bax and Bcl2. The

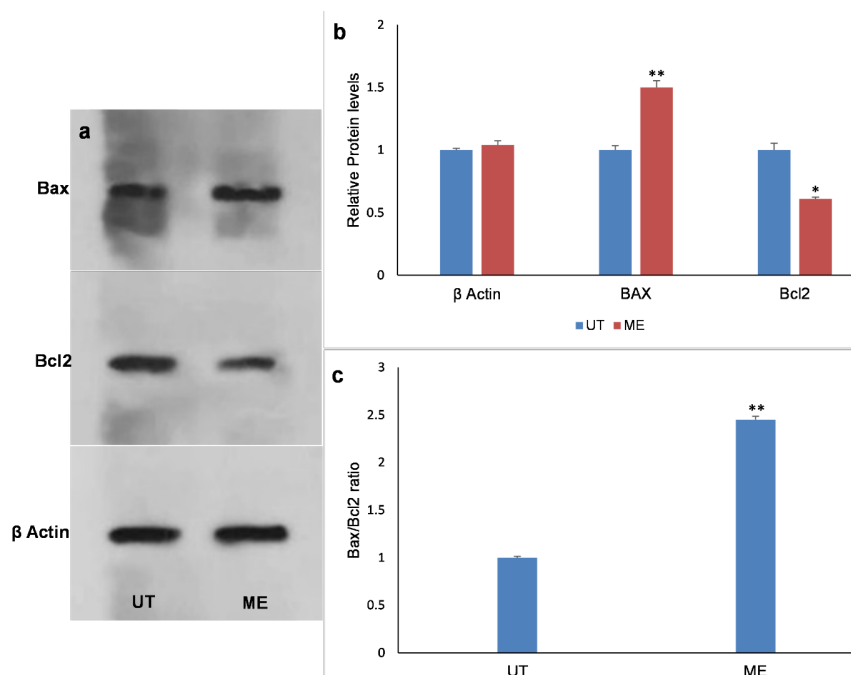


Fig. 8. Western blot analysis of *Bcl2* and *Bax* protein expression in SW480 cells after treatment by ME. (a) *Bax*, *Bcl2*, and β -actin protein bands. (b) The relative protein expression of *Bcl2* and *Bax* compared with that in the control group (UT). (c) The *Bax*: *Bcl2* ratio. Data were expressed as mean values of independent triplicates (mean \pm SD). Indicates a significant difference compared to the control group (* P < 0.05 and ** P < 0.01). ME: methanolic extract, UT: untreated control cells.

Bax/Bcl2 ratio can behave as a rheostat that regulates cell sensitivity to apoptosis. Decreased levels of this ratio may result in cancer cells' resistance to apoptosis. Hence, the *Bax/Bcl2* ratio can impact tumor progression and aggressiveness.⁷⁰⁻⁷² Our findings indicate that *C. turgidus* extract similarly modulates apoptotic pathways, as evidenced by increased *Bax* expression and reduced *Bcl2* levels. Given the established role of *Bax/Bcl2* as a key determinant of apoptotic sensitivity,⁷³ these results suggest that *C. turgidus* metabolites may act via the mitochondrial apoptotic pathway. To improve the clarity and robustness of our apoptosis assay, we have refined our Annexin V flow cytometry data presentation by explicitly reporting the percentages of early apoptosis, late apoptosis, and necrosis. This adjustment ensures a more precise interpretation of apoptotic responses following treatment with *C. turgidus* extract.

Beyond apoptosis induction, drug delivery remains a critical aspect of CRC therapy. Given the increasing use of nanoformulations to enhance the stability and bioavailability of natural compounds,⁷⁴ metabolites of *C. turgidus* can be formulated and used as nanosized drug delivery systems. Lipid-based carriers, polymeric nanoparticles, and liposomal formulations have demonstrated the ability to improve the pharmacokinetics and targeted delivery of marine bioactive compounds.^{75,76} Although our study did not specifically investigate delivery strategies, these insights provide a foundation for future formulation-based studies.

The bioactivity of *C. turgidus* extract is likely attributable to different types of metabolites, including (i) lipopeptides (e.g., cyclic depsipeptides micropeptins, and linear peptides aeruginosins) that interfere with proteolytic enzymes and cellular signaling pathways and used in inflammatory diseases and certain types of cancer, (ii) polyketides (e.g., curacin, and apratoxin) that inhibit microtubule polymerization with potential therapeutic applications in oncology, (iii) indole alkaloids, which disrupt DNA replication and interfere with cell cycle progression, (iv) cyclic peptide toxins (e.g., microcystins, and nodularins) that inhibit protein phosphatases and dysregulate oncogenic pathways, (v) mycosporine-like amino acids, which exhibit photoprotective and antioxidative properties.

Although our study provides compelling evidence supporting the anticancer potential of *C. turgidus*, additional studies are warranted to elucidate the precise molecular interactions underlying its cytotoxic activity. Future research should focus on metabolomic profiling, pharmacokinetic studies, and in vivo validation to further explore the therapeutic applicability of *C. turgidus* bioactive compounds in CRC treatment.

Conclusion

Microalgae and cyanobacteria are recognized as crucial organisms across diverse ecosystems, largely due to their varied structures and capacity to synthesize a wide range of bioactive compounds. While extensive information exists

regarding the compounds produced by microalgae and their biological properties, ongoing research continues to explore their potential applications in drug development and the production of innovative industrial materials. Consequently, there is a pressing need to investigate the therapeutic potentials and applications of various species of microalgae and cyanobacteria. A recent study focusing on the ME from the *C. turgidus* strain KANI revealed significant anticancer activity in vitro. This activity may be positively correlated with the presence of bioactive compounds such as fatty acids, polyphenolic compounds, and anthocyanins. These metabolites are well-known for their antioxidant properties, which allow them to combat free radical oxidation, exhibit therapeutic benefits, and perform essential physiological functions. This research offers valuable insights into the in vitro effects of the ME from *C. turgidus* strain KANI. In summary, the identification and characterization of these metabolites hold great promise for the development of new antitumor agents and could pave the way for innovative approaches to cancer therapy in the future.

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Competing Interests

The authors declare that the research was carried out without any commercial or financial relationships that could be perceived as a potential conflict of interest.

Ethical Approval

This is an in vitro exploratory study. The Institutional Research Ethics Committee has confirmed that no ethical approval is required.

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Research Highlights

What is the current knowledge?

- Microalgae and cyanobacteria produce bioactive compounds with antioxidant and anticancer effects.
- *Chroococcus turgidus* has demonstrated antioxidant, anti-inflammatory, antibacterial, and antiviral properties.
- The Bax/Bcl2 pathway plays a key role in apoptosis induction in colorectal cancer cells.

What is new here?

- *Chroococcus turgidus* methanolic extract inhibits colorectal cancer cell proliferation in a dose- and time-dependent manner.
- Bioactive metabolites, including phenols, flavonoids, and fatty acids, contribute to its anticancer effects.
- Apoptosis induction is confirmed by nuclear fragmentation, Annexin V staining, and Bax/Bcl2 regulation.

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