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Heliotropium ramosissimum metabolic profiling, in silico and in vitro evaluation with potent selective cytotoxicity against colorectal carcinoma

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Heliotropium is a genus of the Boraginaceae family. Its members are used in many traditional and folklore medicines to treat several ailments. Despite this widespread usage, only a few evidencebased scientific studies investigated and identified its phytoconstituents. Herein, we documented the chemical profile of the Heliotropium ramosissimum methanolic extract using gas chromatographymass spectrometry (GC-MS) and liquid chromatography-tandem mass spectrometry (LC-ESI-MS/ MS) and assessed its antioxidant and cytotoxic effects. The methanolic extract exhibited high phenolic content (179.74 ± 0.58 μg/mL) and high flavonoid content (53.18 ± 0.60 μg/mL). The GC-MS analysis of the lipoidal matter allowed us to identify 41 compounds with high percentages of 1,2-benzenedicarboxylic acid, bis(2-methoxyethyl) ester (23.91%), and 6,10,14-trimethylpentadecan-2-one (18.74%). Thirty-two phytomolecules were tentatively identified from the methanolic extract of H. ramosissimum using LC-MS/MS. These compounds belonged to several phytochemical classes such as phenolic acids, alkaloids, coumarins, and flavonoids. Furthermore, we assessed the antioxidant activity of the methanolic extract by DPPH assay and oxygen radical absorbance capacity assay, which yielded IC_{so} values of 414.30 μg/mL and 170.03 ± 44.40 μM TE/equivalent, respectively. We also assessed the cytotoxicity of the methanolic extract on seven different cell lines; Colo-205, A-375, HeLa, HepG-2, H-460, and OEC showed that it selectively killed cancer cells with particularly potent cytotoxicity against Colo-205 without affecting normal cells. Further studies revealed that the extract induced apoptosis and/or necrosis on Colo-205 cell line at an IC_{s0} of 18.60 μ g/mL. Finally, we conducted molecular docking on the LC-ESI-MS/MS-identified compounds against colon cancer antigen 10 to find potentially cytotoxic compounds. Binding score energy analysis showed that isochlorogenic acid and orientin had the highest affinity for the colon cancer antigen 10 protein, with binding scores of (-13.2001) and (-13.5655) kcal/mol, respectively. These findings suggest that Heliotropium ramosissimum contains potent therapeutic candidates for colorectal cancer treatment.

Abbreviations

A-375 Human melanoma cell line
AlCl₃ Aluminum chloride
Colo-205 Colorectal cancer cell line
DPPH 2,2-Diphenyl-1-picrylhydrazyl
GAE Gallic acid equivalents

GC-MS Gas chromatography-mass spectrometry

H-460 Large cell lung cancer cell line HeLa Cervical cancer cell line

HepG-2 Hepatocellular carcinoma cell line

LC-ESI-MS/MS Liquid chromatography-electrospray ionization-mass spectrometry

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OEC Oral epithelial cell line

ORAC Oxygen radical absorbance capacity

RE Rutin equivalents SRB Sulforhodamine B TE Trolox equivalent

Family Boraginaceae is one of the largest families that include hundreds of genera and nearly 2 thousand species. Members of this family can be found in both tropical and temperate climates, especially in the Mediterranean. *Heliotropium*, a genus that belongs to this family, contains perennial and annual herbs with scorpioid cymes in their inflorescences as well as the morphology of the greatly modified stigmatic head in the flower¹. In various traditional and folklore medical systems, *Heliotropium* species have been used to treat several illnesses. Various parts of the plant, including the leaves, roots, flowers, seeds, and the whole plant, are used to treat a variety of ailments². In folk medicine, *H. aegyptiacum* is used to treat snake bites and scorpion stings³, while *H. indicum* is reported to be used against infected gums⁴, head lice⁵, rheumatism⁶, and gonorrhea⁷. *H. strigosum* has laxative, diuretic effects⁸, as well as its activity against breast abscesses⁹. *H. crispum* is used as a cooling agent and exerted a lactagogue effect in cattle¹⁰. *H. steudneri* is used to stop bleeding and to prevent infection⁴, while *H. ramosissimum* is used to treat burns⁴. Gout, inflammation, menstrual dysfunction, skin disorders, noxious bites, and rheumatism have all been confirmed to be treated with *Heliotropium* species².

Heliotropium species contain phytochemicals that are bioactive and have powerful healing properties. Several classes of organic compounds viz. flavonoids, pyrrolizidine alkaloids, terpenoids, and quinones are very plentifully present in the Heliotropium genus¹¹. Afghanistan, Burkina Faso, Canary Islands, Cape Verde, Chad, Djibouti, Egypt, Gulf States, Iran, Iraq, Lebanon-Syria, Libya, Mauritania, Niger, Saudi Arabia, Senegal, Sinai, Sudan, Turkmenistan, and Yemen are all home to Heliotropium ramosissimum (Lehm.) DC¹². Few studies concerning the isolation of phytoconstituents from this species were reported¹³, in addition to evaluation of its total phenolic content and total antioxidant capacity using different assays¹⁴. This scarcity of knowledge on its phytochemical and biological properties encouraged us to document its phytochemical properties, antioxidant activity, and in vitro cytotoxic activity toward various cell lines. Moreover, we explored the compounds potentially responsible for the methanolic extract's cytotoxic activity through *in-silico* studies.

Materials and methods

Statement. All experiments and methods including collection of the plant were performed following relevant national, and international guidelines and legislation of the Faculty of Pharmacy, University of Sadat City, Sadat City, Egypt.

Plant material. We collected *Heliotropium ramosissimum* (Lehm.) DC. flowering aerial parts (Fig. 1) in March 2020 from the El-Sadat City desert region, Sadat City, Egypt with license approval from the Faculty of Pharmacy, University of Sadat City, Sadat City, Egypt according to relevant guidelines and regulations. The plant material was kindly identified by Prof. Dr. A. A. Fayed, Professor of Plant Taxonomy, Faculty of Science, Assiut University, Assiut, Egypt. We deposited a voucher sample (alphabetically ordered under the letter "H") in the Herbarium of the Faculty of Science, Assiut University, Assiut, Egypt.

Extraction and fractionation. The air-dried powdered aerial parts of H. ramosissimum (Lehm.) DC. (1 kg) were defatted with n-hexane (3×1.5 L) to prepare the lipoidal matter and then extracted with methanol (3×1.5 L) to prepare the total methanolic extract used in this study¹⁵. We performed the extractions in each solvent until exhaustion. After completing the process, we removed each solvent under reduced pressure using a rotary evaporator (Acculab, USA) at 50 °C. The n-hexane yielded 5 g of residue, and the total methanolic extracts weighed 10 g. We stored the extracts in a vacuum desiccator until further use^{16,17}.

Phytochemical studies. *Phytochemical screening.* We identified the presence of phytochemical classes in freshly prepared crude extracts of the flowering aerial parts of *H. ramosissimum* using standard colorimetric procedures ^{18,19}.

Estimation of the total phenolic and flavonoid contents. The total phenolic content was calculated as gallic acid equivalents (GAE) per g of the sample using the Folin–Ciocalteu reagent and a calibration curve prepared with gallic acid²⁰. Moreover, the total flavonoid content was determined as rutin equivalents (RE) per g of the sample using aluminum chloride (AlCl₃) colorimetric assay²⁰.

Metabolomic analysis. *GC–MS analysis of the lipoidal matter.* The chemical composition of the lipoidal matter of the aerial parts of *H. ramosissimum* was determined using a Trace GC-TSQ mass spectrometer (Thermo Fisher Scientific, Austin, TX, USA) with a direct capillary column TG–5MS (Thermo Fisher Scientific, Austin, TX, USA) (30 m × 0.25 mm × 0.25 μm film thickness). The initial column oven temperature was 50 °C. It was then increased to 250 °C at 5 °C/min, held for 2 min, increased to the final temperature of 300 °C at 30 °C/min, and held for 2 min. The injector and MS transfer line were kept at 270 °C and 260 °C, respectively. The carrier gas (helium) had a constant flow rate of 1 mL/min. The solvent delay was 4 min, and diluted 1 μL samples were injected automatically using Autosampler AS1300 coupled with GC in split mode. We collected electron ionization mass spectra at an ionization voltage of 70 eV over the *m/z* range 50–650 in full scan mode. We set the



Figure 1. A Photo of the *H. ramosissimum* (Lehm.) DC. flowering aerial from the place of collection (taken by the corresponding author in March 2020 from El-Sadat City desert region, Sadat City, Egypt).

ion source temperature to 200 °C. We identified the components by comparing their mass spectra with those of the WILEY 09 and NIST 14 mass spectral libraries 16.

LC-ESI-MS/ MS profiling. We performed the LC-ESI-MS/MS analysis of the methanolic extract on an ExionLC AC system coupled with a SCIEX Triple Quad 5500 + MS/MS system equipped with an electrospray ionization (ESI) system. The samples were eluted on an Ascentis C18 Column (4.6×150 mm, 3 μm). Mobile phases consisted of eluent A (0.1% formic acid) and eluent B (acetonitrile, LC grade). The mobile phase gradient was programmed as follows: 10% B at 0-1 min, 10%-90% B at 1-21 min, 90% B at 21-25 min, and 10% at 25.01-28 min. The flow rate was 0.5 mL/min, and the injection volume was 10 μL. MS/MS analysis used positive and negative ionization modes with a scan (EMS-IDA-EPI) from 100 to 1000 Da for MS1 with the following parameters: curtain gas, 25 psi; Ion Spray voltage, 5500 and -4500 v for positive and negative modes, respectively; source temperature, 500 °C; ion source gas 1% 2, 45% psi and from 50% to 800% Da for MS2; declustering potential, 80%; collision energy, 35% and -35% for positive and negative modes, respectively; and collision energy spread, 20^{21} . We identified the compounds using MS-DIAL software version 4.70.

Free radical scavenging activity assessment by DPPH assay. We prepared 1000 and 100 µg/mL solutions from the methanolic extract to identify a range within which the inhibitory concentration 50 (IC $_{50}$) lay. We serially diluted the solutions exceeding 50% five times. We prepared a 100 µM Trolox stock solution in methanol. From this stock solution, we prepared 50, 40, 30, 20, 15, 10, and 5 µM solutions. We performed the 2,2-diphenyl1-picryl-hydrazyl-hydrate (DPPH) free radical assay as described by Boly et al., 2016^{17,20}. Briefly, we added 100 µL of freshly prepared DPPH reagent (0.1% in methanol) to 100 µL of the sample in a 96-well plate (n = 6) that we incubated at room temperature for 30 min in the dark. Next, we measured the reduction in DPPH color intensity at 540 nm using the microplate reader FluoStar Omega. We presented the data as mean ± standard deviation according to the following equation:

$$Percentage\ inhibition = \frac{Average\ absorbance\ of\ blank-Average\ absorbance\ of\ the\ test}{Average\ absorbance\ of\ blank} \times 100.$$

Assessment of the oxygen radical absorbance capacity (ORAC assay). We prepared a 1 mM Trolox stock solution and performed nine serial dilutions to obtain 400, 300, 200, 150, 100, 75, 50, 25, and 12.5 μ M solutions. We prepared a 400 μ g/mL sample solution in MeOH. We performed the assay as described by Liang et al., 2014²², with minor modifications. Briefly, we incubated 12.5 μ L of the prepared sample(s) with 75 μ L of fluorescein (10 nM) for 30 min at 37 °C. Next, we measured background fluorescence (485 nm excitation and 520 nm emission) for three cycles (cycle time: 90 s). Then, we immediately added 12.5 μ L of freshly prepared 2,2'-azobis(2-amidinopropane) (240 mM) to each well and continuously measured fluorescence (485 nm excitation and 520 nm emission) for 2.5 h (100 cycles of 90 s each). We presented the data as mean (n=3) \pm standard deviation and calculated the antioxidant effect of the extract as μ M Trolox equivalents by substitution in the linear regression equation: Y = 32,356.3X + 989,769.9 ($R^2=0.9957$).

Assessment of cytotoxic activity. *Cell culture.* We obtained colorectal cancer (Colo-205), human melanoma (A-375), cervical cancer (HeLa), hepatocellular carcinoma (Hep G-2), large cell lung cancer (H-460), and oral epithelial (OEC) cells from Nawah Scientific Inc. (Mokatam, Cairo, Egypt). We maintained them in RPMI medium supplemented with 100 mg/mL of streptomycin, 100 units/mL of penicillin, and 10% of heatinactivated fetal bovine serum and incubated them in humidified 5% (v/v) CO_2 atmosphere at 37 ° C^{23} .

Cytotoxicity assay. We assessed cell viability through a sulforhodamine B (SRB) assay. We added 100 μ L of cell suspension (5×10³ cells) to 96-well plates and incubated them in a complete medium for 24 h. We then treated the cells with 100 μ L of medium containing samples at different concentrations (10 and 100 μ g/mL). After 72 h of exposure, we fixed the cells by replacing the medium with 150 μ L of 10% trichloroacetic acid and incubated them at 4 °C for 1 h. Next, we removed the trichloroacetic acid solution and washed the cells five times with distilled water. We then added 70 μ L of SRB solution (0.4% w/v) and incubated the mixture in a dark place at room temperature for 10 min. We washed the plates three times with 1% acetic acid and allowed them to air-dry overnight. Then, we added 150 μ L of TRIS (10 mM) to dissolve the protein-bound SRB stain and measured the absorbance at 540 nm using a BMG LABTECH-FLUOstar Omega microplate reader (Ortenberg, Germany)²³.

Annexin-based apoptosis assay. Colo-205 cells were treated with either doxorubicin (10 μ M) as a positive control or the total methanolic extract of *H. ramosissimum* for 48 h. Next, we collected the cells (10⁵ cells) by trypsinization and washed them twice with ice-cold phosphate-buffered saline (PBS, pH 7.4). We then incubated the cells in the dark with 0.5 mL of Annexin V-FITC/propidium iodide (PI) solution for 30 min at room temperature according to the manufacturer's protocol (Annexin V-PI staining apoptosis detection kit from Abcam Inc., Cambridge Science Park, Cambridge, UK). After staining, we injected the cells into an ACEA Novocyte flow cytometer (ACEA Biosciences Inc., San Diego, CA, USA) and detected the FITC and PI fluorescent signals using FL1 and FL2 signal detectors, respectively ($\lambda_{\text{ex/em}} = 488/530$ nm for FITC and $\lambda_{\text{ex/em}} = 535/617$ nm for PI). For each sample, we acquired 12,000 events and quantified the FITC- and PI-positive cells by quadrant analysis using ACEA NovoExpress software (ACEA Biosciences Inc., San Diego, CA, USA)²⁴.

Cell cycle distribution analysis. After treating the cells (10^5 cells) with the total methanolic extract of H. ramosissimum for 48 h or paclitaxel ($1~\mu M$) for 24 h, as a positive control, we collected them by trypsinization and washed them twice with ice-cold PBS (pH 7.4). We then resuspended the cells in 2 mL of 60% ice-cold ethanol and incubated them at 4 °C for 1 h for fixation. Next, we washed the fixed cells twice with PBS (pH 7.4) and resuspended them in 1 mL of PBS containing 50 $\mu g/mL$ RNAase A and 10 $\mu g/mL$ PI. After 20 min of incubation in the dark at 37 °C, we analyzed the cells' DNA contents by flow cytometry using an FL2 ($\lambda_{ex/em}$ 535/617 nm) signal detector (ACEA Novocyte flow cytometer, ACEA Biosciences Inc., San Diego, CA, USA). For each sample, 12,000 events were acquired. We determined cell cycle distribution using ACEA NovoExpress software (ACEA Biosciences Inc., San Diego, CA, USA).

Statistical analysis. We carried out triplicate experiments and analyzed data using *Microsoft Excel*. We determined the IC_{50} using *GraphPad Prism 5* by converting the concentrations to their logarithmic value and using the "non-linear inhibitor regression equation (log (inhibitor) vs. normalized response–variable slope equation)" function²⁶.

Molecular docking. We performed docking studies using Molecular Operating Environment (MOE, 2014.0901)²⁷. We retrieved the three-dimensional (3D) structures of serologically defined colon cancer antigen 10 (PDB ID: 2HQ6) determined by X-ray diffraction (resolution: 1.75 Å) from the RCSB Protein Data Bank (https://www.rcsb.org/) and used it as a target in our molecular docking experiments²⁸. We prepared and optimized the structure of the receptor protein using MOE's ligx function with default settings. We drew the structures of the identified compounds in Chemdarw 17.0.0.206 in MOL format and performed 3D protonation, partial energy correction, and energy minimization using Merck molecular force field (MMFF94x). We determined the receptor active sites using MOE's site finder tool. We carried out the molecular docking analysis to predict the receptor–ligand interaction using flexible ligand-fixed receptor docking parameters with Triangle Matcher placement (scoring: London dG; retain: 30) and force field refinement (rescoring: London dG; retain: 10). We selected the most stable protein–ligand interactions based on their S-score minimum energy and root mean square deviation (RMSD). We recorded the docking score, RSMD, and 2D and 3D interactions^{29,30}.

Results and discussion

Phytochemical screening. We carried out phytochemical screening on *H. ramosissimum* to identify the different chemical classes of the total methanolic extract active constituents using different reagents. The preliminary screening revealed the presence of alkaloids, terpenoids, saponins, tannins, flavonoids, coumarins, polyphenolics, and reducing sugars.

Total phenolic and flavonoid contents. We estimated the total phenolic content in the methanolic extract of H. ramosissimum to be 179.74 ± 0.58 µg/mL, in gallic acid equivalents. The equation for the standard curve was Y = 0.0031x - 0.0564 (Fig. S1), with $R^2 = 0.9961$. Moreover, we measured a total flavonoid content of 53.18 ± 0.60 µg/mL, in rutin equivalents. The equation for the standard curve was Y = 0.0032x + 0.0398 (Fig. S2), with $R^2 = 0.9981$.

Metabolomic analysis. *GC–MS analysis of the lipoidal matter.* We performed a GC–MS analysis of the plant's lipoidal matter to address the lack of information regarding its phytoconstituents. The GC–MS chromatogram (Fig. S3) of the lipoidal matter of *H. ramosissimum* revealed the presence of 41 compounds, and we identified 30 (91.73%) of them by comparing their mass spectra data (retention time and peak area) with the Wiley spectral library. We left 11 (8.25%) compounds unidentified. The most abundant compound was 1, 2-benzenedi-

Peak no	R _t (min)	Relative (%)	Mol. formula	[M+] m/z	Identified Compounds*	References
1	15.14	0.65	C ₁₇ H ₃₆	240	2,6,10-Trimethyltetradecane	31
2	16.72	0.79	C ₁₃ H ₁₈ O ₂	206	3,4-Dihydro-2H-1,5-(3"-t-butyl) benzodioxepine	32
3	18.23	0.35	C ₁₈ H ₃₄ O ₂	282	Trans-13-octadecenoic acid	33
4	18.61	0.70	C ₈ H ₆ N ₄ O ₅	238	2,4-Imidazolidinedione, 1-[[(5-nitro-2-furanyl) methylene]amino]-	34
5	19.86	0.39	C ₁₆ H ₃₀ O ₄ Si3	370	Benzoic acid, 2,4-bis[(trimethylsilyl)oxy]-, trimethylsilyl ester	35
6	20.10	0.40	C ₁₇ H ₃₆ O	256	2-Methylhexadecan-1-ol	36
7	21.27	0.62	$C_{20}H_{38}O_2$	310	cis-13-Eicosenoic acid	37
8	22.61	0.45	C ₁₉ H ₃₆ O ₂	296	cis-10-Nonadecenoic acid	38
9	22.74	0.55	C ₃₂ H ₆₆	450	Dotriacontane	39
10	23.25	0.21	C ₁₈ H ₁₆ O ₇	344	4H-1-Benzopyran-4-one, 2-(3,4-dimethoxyphenyl)- 3,5-dihydroxy-7-methoxy	40
11	23.73	18.74	C ₁₈ H ₃₆ O	268	6,10,14-Trimethylpentadecan-2-one	41
12	24.27	0.86	C ₂₆ H ₄₂ O ₄	418	1,2-Benzenedicarboxylic acid, tetradecyl ester	42
13	24.65	0.42	C ₂₁ H ₃₆	288	14-α-Pregnane	43
14	24.76	0.22	C ₁₈ H ₃₄ O	266	11-Octadecenal	44
15	25.23	0.54	$C_{19}H_{22}O_6$	346	Isochiapin B	45
16	22.35	11.83	$C_{17}H_{34}O_2$	270	Hexadecanoic acid methyl ester	46
17	26.15	23.91	C ₁₄ H ₁₈ O ₆	282	1,2-Benzenedicarboxylic acid, bis (2-methoxyethyl) ester	47
18	26.66	13.68	C ₁₈ H ₃₆ O ₂	284	Ethyl hexadecanoate	48
19	27.28	0.51	$C_{18}H_{36}O_2$	284	Methyl 14-methylheptadecanoate	49
20	28.54	2.51	$C_{19}H_{34}O_2$	294	Linolelaidic acid, methyl ester	50
21	28.65	1.69	C ₂₂ H ₃₈ O ₂	334	Methyl 8-[2-((2-[(2-ethylcyclopropyl) methyl] cyclopropy) methyl) cyclopropyl] octanoate	51
22	29.12	1.78	$C_{19}H_{38}O_2$	298	Methyl octadecanoate	52
23	29.73	1.26	$C_{20}H_{36}O_2$	308	Linoleic acid ethyl ester	48
24	29.84	1.02	$C_{19}H_{34}O_2$	294	(3E,12Z)-1,3,12-Nonadecatriene-5,14-diol	53
25	30.15	0.27	$C_{19}H_{38}O_4$	330	2,3-Dihydroxypropyl palmitate	54
26	30.32	1.74	$C_{20}H_{40}O_2$	312	Ethyl octadecanoate	55
27	31.59	0.51			Unknown	-
28	32.05	0.45	C ₁₈ H ₃₀ O ₆₀	268	2,2,3,3,4,4 Hexadeutero octadecanal	56
29	33.07	1.47			Unknown	-
30	33.68	0.46	$C_{21}H_{40}O_2$	324	4,8,12,16-Tetramethylheptadecan-4-olide	57
31	33.94	0.45			Unknown	-
32	35.28	0.45			Unknown	-
33	36.11	2.42	$C_{24}H_{38}O_4$	390	1,2-Benzenedicarboxylic acid, dioctyl ester	58
34	36.80	0.32			Unknown	-
35	37.91	0.79			Unknown -	
36	38.65	0.48			Unknown -	
37	38.82	0.42			Unknown -	
38	39.01	1.66			Unknown -	
39	39.14	2.31	$C_{29}H_{50}O_4$	462	α-Tocospiro A 58	
40	39.23	1.49			Unknown	-
41	39.58	0.21			Unknown	_

Table 1. GC–MS data of the n-hexane fraction of H. ramosissimum (Lehm.) DC. aerial parts. * The identification of n-hexane constituents based on the comparison of the mass spectral data with those of Mass Spectral Library (2011) and Wiley Registry of Mass Spectral Data 8th edition and literature.

carboxylic acid, bis (2-methoxyethyl) ester (23.91%), followed by 6,10,14-trimethylpentadecan-2-one (18.74%), ethyl hexadecanoate (13.68%), and hexadecanoic acid methyl ester (11.83%) (Table 1 and Fig. S4).

LC–ESI–MS/MS profiling. We identified the compounds of the H. ramosissimum methanolic extract by LC–MS/MS. We compared their retention time (R_t), mass, and MS2 with standards-reported literature, and databases (TMIC and MassBank). In total, we tentatively identified 32 compounds, including alkaloids, flavonoids, coumarins, phenolic acids, and their derivatives. We identified 17 compounds in negative mode and 15 in posi-

No	R _t (min)	[M+H]+	Fragments	Identified Compound	Chemical Class	Ref
1	0.42	149	79, 105, 131	Cinnamic acid	Phenolic acids	59
2	9.28	191	115, 119, 147, 148	7-Methoxy-4-methylcoumarin	Coumarins	59
3	10.36	291	273, 139, 123	(epi)-Catechin	flavan -3-ol	60
4	10.94	300	138, 139, 157	Lycopsamine	Alkaloids	61-63
5	11.22	314	138, 156, 269	Heliospathine	Alkaloids	61,62
6	13.20	286	153, 127, 109	Trachelanthamine	Alkaloids	64
7	15.70	185	126, 80	Methyl gallate	Phenolics	65
8	15.95	277	134, 175, 217, 241	Stearidonic acid	Fatty acids	66
9	18.09	289	153, 127, 109	Eriodictyol	Flavonoids	67
10	19.57	287	241, 261, 213, 153	Kaempferol	Flavonoids	68
11	19.85	305	287, 259	Taxifolin	Flavonoids	68
12	19.89	330	121, 139, 241	Europine	Alkaloids	61,69,70
13	22.72	398	107, 121, 139	Heliosupine	Alkaloids	61,71,72
14	25.80	303	285, 275, 257, 229	Quercetin	Flavonoids	68
15	27.15	156	81, 112, 139	Retronecine	Alkaloids	61,63,73,74

Table 2. Metabolites identified in the methanolic extract of *H. ramosissimum* (Lehm.) DC. using positive mode LC–ESI–MS/MS.

No	R _t (min)	[M-H] ⁻	Fragments	Identified Compound	Chemical Class	Ref
1	2.21	115	97, 71	Maleic acid	Dicarboxylic acids	59,75
2	2.59	117	73, 98, , 116	Succinic acid	Dicarboxylic acids	76
3	4.14	181	125, 137, 153	2-Ethoxy-4,5-dihydroxybenzaldehyde	Phenolics	77
4	4.18	241	105, 195, 225	6-Deoxycochinolide	Benzofuran	78
5	4.48	175	113, 115, 131, 157	2-Isopropylmalic acid	Organic acids	59,79
6	4.61	138	79, 107, 110	7-Hydroxy-1-methylenepyrrolizidine	Alkaloids	80
7	4.70	153	153, 135, 123, 109	Protocatechuic acid	Phenolic acids	59,81
8	5.55	137	93, 109, 119	4-Hydroxybenzoic acid	Phenolic acids	59,81,82
9	5.6	179	135, 143, 161, 107	Caffeic acid	Phenolic acids	83
10	9.39	177	105, 133, 146, 176	4-Methoxycinnamic acid	Phenolic acids	59
11	10.01	289	243, 245	Filifolinoic acid	Phenolic acids	84,85
12	10.77	161	105, 117, 133	Umbelliferone	Coumarins	59,86
13	16.92	515	515, 353, 179, 173	4,5-di-O-caffeoylquinic acid (isochlorogenic acid)	Phenolic acids	87
14	20.05	337	337, 191, 163	3-O-p-coumaryl quinic acid	Phenolic acids	87,88
15	24.9	447	313, 319, 327, 357	Luteolin 8-C-Glucoside (Orientin)	Flavonoids	89
16	26.73	367	367, 191	3-O-feruloyl quinic acid	Phenolic acids	90
17	27.89	233	109, 121, 145, 175	Tournefolin C	Flavonoids	77

Table 3. Metabolites identified in the methanolic extract of *H. ramosissimum* (Lehm.) DC. using negative mode LC–ESI–MS/ MS.

tive mode. Tables 2 and 3 list all identified compounds ordered by relative retention times and their structures were supplied (Figs. S5, S6, S74, S8).

Free radical scavenging activity. The DPPH spectrophotometric assay is one of the most reliable and widely used methods to estimate the free radical scavenging effect of different plant extracts. The percent of inhibition values in the initial screening step of the two concentrations, namely, 1000 and 100 µg/mL, were 80.26 ± 1.50 ($^{>}50$) and 6.50 ± 0.73 ($^{<}50$), respectively. We serially diluted the extract that exceeded 50% inhibition (1000 µg/mL) to provide five concentrations that we tested to determine that the IC50 was 414.30 µg/mL (Fig. S9). For reference, Trolox has an IC50 of 24.42 ±0.87 µM. The H. ramosissimum methanolic extract showed a potent radical-scavenging effect, which may be attributed to its total phenolic and flavonoid contents.

Assessment of ORAC assay. We evaluated the antioxidant activity of the H. ramosissimum methanolic extract using the ORAC assay (Fig. S10). The antioxidant activity of the extract (Fig. 2) was $170.03\pm44.45~\mu M$ TE/equivalent, which is higher than that of Trolox.

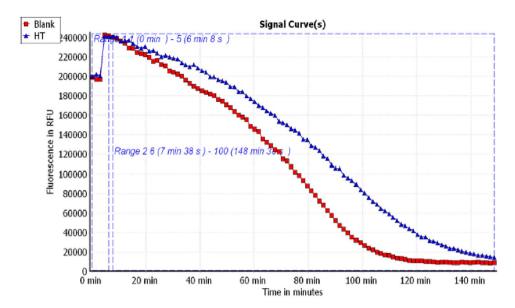


Figure 2. Signal curves of total methanolic extract of *H. ramosissimum* (Lehm.) DC. aerial parts (HT, blue color) and blank indicating the decay of fluoresceine upon applying the extract.

	Cell viability %								
	Cancer cell li	Normal cell line							
Conc. of the tested sample	Colo-205	A-375	HeLa	HepG-2	H-460	MCF-7	OEC		
10 μg/mL	75.95 ± 0.81	98.51 ± 0.44	98.03 ± 1.44	99.60 ± 0.19	95.93 ± 1.34	96.51 ± 0.73	98.94±0.77		
100 μg/mL	2.24 ± 0.37	3.65 ± 0.93	9.46±0.37	23.82 ± 1.50	42.85 ± 0.26	61.66 ± 1.32	51.62 ± 0.48		

Table 4. Cytotoxicity SRB quick screening results of the total methanolic extract of *H. ramosissimum* (Lehm.) DC. aerial parts. \pm SD (n = 3).

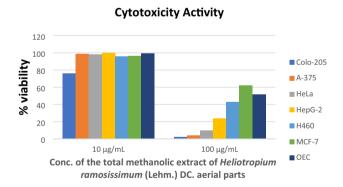


Figure 3. SRB cytotoxicity quick screening results of the total methanolic extract of *H. ramosissimum* (Lehm.) DC. aerial parts on different cell lines.

Cytotoxicity assay. We screened the cytotoxicity of the H. ramosissimum methanolic extract at 10 and 100 µg/mL on various cell lines (Table 4 and Fig. 3). We performed an SRB quick screening assay on six cancer cell lines, namely, colorectal cancer (Colo-205), human melanoma (A-375), cervical cancer (HeLa), hepatocellular carcinoma (HepG-2), large cell lung cancer (H-460) cells, and a normal cell line, that is, oral epithelial cells (OEC). The H. ramosissimum methanolic extract exhibited potent activity on all tested cell lines. The most sensitive cell line was Colo-205, followed by A-375, HeLa, and H-460, and the least sensitive was the MCF-7 cell line. Because the H. ramosissimum methanolic extract showed the highest activity against colorectal carcinoma (Colo-205) cells—in excellent agreement with the in silico study—we further investigated its IC_{50} and mechanism of action through cell cycle and apoptosis flow cytometry assays on this cell line.

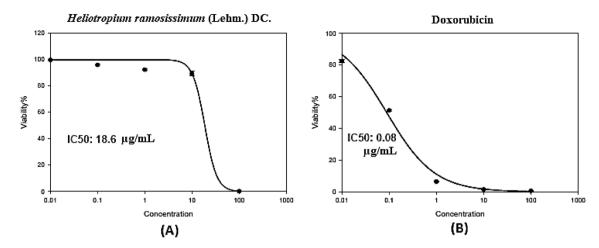


Figure 4. *In-vitro* SRB cytotoxicity assay of (**A**) the total methanolic extract of *H. ramosissimum* (Lehm.) DC. aerial parts; (**B**) Doxorubicin; in increasing concentrations $(0.01-100 \,\mu\text{g/mL})$ against colorectal carcinoma (Colo-205) cell lines. Data points are expressed as mean \pm SD (n=3).

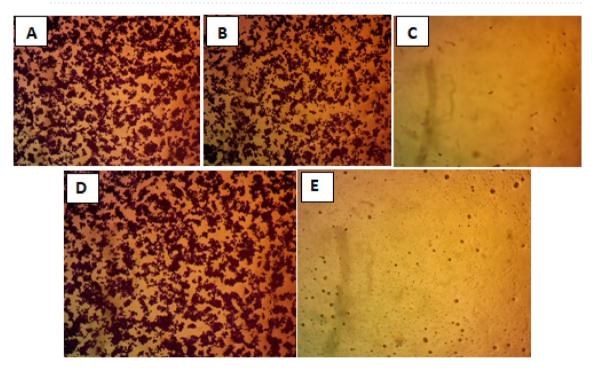


Figure 5. Optical microscope stained images of SRB cytotoxicity assay against Colo-205 cell line (**A**) Negative control, (**B**) Doxorobucin (0.01 μg/mL), (**C**) Doxorobucin (100 μg/mL), (**D**) the total methanolic extract of *H. ramosissimum* (Lehm.) DC. aerial parts (0.01 μg/mL), (**E**) the total methanolic extract of *H. ramosissimum* (Lehm.) DC. aerial parts (100 μg/mL), magnification power: X 100.

We assessed cell viability at five different concentrations (0.01, 0.1, 1, 10, and 100 μ g/mL) and used doxorubicin as a standard cytotoxic drug. The SRB assay (Figs. 4, 5) revealed that the *H. ramosissimum* methanolic extract possesses a dose-dependent cytotoxic effect with an IC $_{50}$ of 18.60 μ g/mL (Fig. 4A) and that of doxorubicin is 0.08 μ g/mL (Fig. 4B). The *H. ramosissimum* extract decreased cell viability from 75.95% \pm 0.81% at 10 μ g/mL to 2.24% \pm 0.37% at 100 μ g/mL (Table 4).

The optical microscope staining images (Fig. 5) show the results of the SRB cytotoxicity assay against Colo-205 cell line of both total Colo-205 cells for two *H. ramosissimum* methanolic extract concentrations (0.01 and 100 μ g/mL), doxorubicin at the same concentrations, and negative control. The figure clearly shows that at 0.01 μ g/mL, neither the extract nor doxorubicin caused significant morphological changes. Meanwhile, significant changes occurred at 100 μ g/mL, confirming the dose-dependent character of the cytotoxicity of the extract. To investigate the safety of the *H. ramosissimum* methanolic extract on normal cells and the selectivity of its cytotoxicity on cancer cells, we performed the cytotoxic activity assay on OECs. The viability of OECs treated with 10 μ g/mL methanolic extract was 98.94% \pm 0.77%. Meanwhile, this concentration showed a potent cytotoxic effect on Colo-205 cells by reducing cell viability to 75.95% \pm 0.81%.

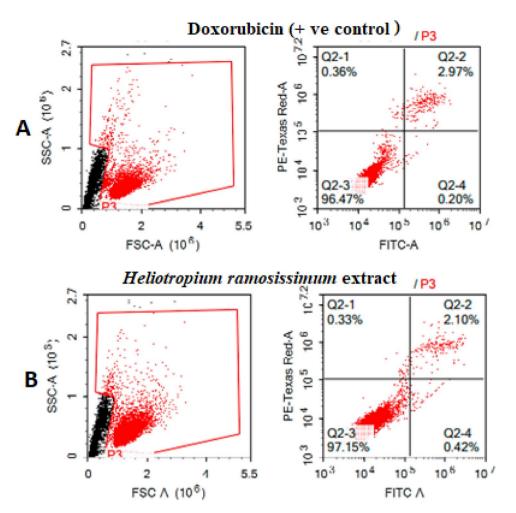


Figure 6. Apoptosis in Colo-205 cells estimated by Annexin V and PI staining followed by flow cytometry analysis. Quadrants represent dead (Q1), late apoptotic (Q2), live (Q3), and (Q4) early apoptotic cells. (**A**) Control, (**B**) the total methanolic extract of *H. ramosissimum* (Lehm.) DC. aerial parts.

To further investigate the mechanism responsible for the cytotoxic activity of the H. ramosissimum methanolic extract, we carried out an apoptosis assay on Colo-205 cells. We used Annexin V-FITC/PI double staining and evaluated the apoptotic rates using flow cytometry. As illustrated in Fig. 6, the apoptotic rate was 56.14% in the positive control group using doxorubicin, whereas treating the cells with 18 μ g/mL of methanolic extract (which is equal to its IC₅₀) significantly increased the apoptotic rate to 71.76%. These results suggest that the extract reduced Colo205 cell viability by inducing apoptosis.

To examine whether the cytotoxicity of the H. ramosissimum methanolic extract on Colo-205 was associated with cell cycle arrest, we conducted a PI-metric cell cycle analysis using flow cytometry (Fig. 7). The cell cycle histograms revealed that cells treated with the extract (18 μ g/mL) had a markedly higher sub-G1 population (63.01%) than those treated with the positive control, paclitaxel (37.33%). This suggests that the extract induced apoptosis and/or necrosis in Colo-205 cells.

We reviewed the literature on the cytotoxic activity of different *Heliotropium* species and found that the n-hexane fraction of the ethanolic extract of H. subulatum aerial parts displayed cytotoxic activity at 3 mg/mL 91 . An in vitro assay against MRC5 human cells revealed that the methanolic extract of H. subulatum aerial parts exerted significant cytotoxicity with an IC $_{50}$ of 13 μ g/mL 92 . Furthermore, the methanolic extract of the dried roots of H. subulatum showed different mortality rates at different concentrations, with a lethal concentration (LC $_{50}$) of 47.86 μ g/mL and LC $_{90}$ of 75.85 μ g/mL in a brine shrimp lethality bioassay 93 . A 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) assay revealed that the whole plant ethanolic extract of H. subulatum had significant antiproliferative activity against SKBR-3 human breast adenocarcinoma cells. The IC $_{50}$ of the extract was 34 ± 9.09 μ g/mL, and that of the standard drug paclitaxel was 22.20 ± 2.30 μ g/mL 94 . These findings suggest that the subulatum methanolic extract exerts a potent cytotoxic effect, especially against colorectal carcinoma, by inducing apoptosis. These results suggest that this extract is a promising candidate for the treatment of this type of cancer.

Molecular docking. In silico molecular docking, techniques have enhanced drug discovery and development by allowing the structure-based exploration of ligand-receptor interactions. The cytotoxicity assays

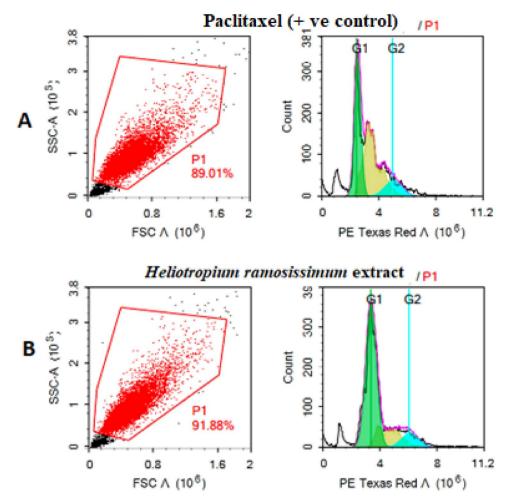


Figure 7. Schematic representation of cell cycle arrest at Sub G1 and G2 phase (**A**) Control, (**B**) the total methanolic extract of *H. ramosissimum* (Lehm.) DC. aerial parts.

revealed that the cytotoxicity of the methanolic extract was more potent against colon cancer than against the other cell lines. We investigated the possible interactions between the 32 phytochemical compounds identified by LC–ESI–MS/MS and the colon cancer antigen 10 binding sites and compared them to the standard anticancer drug doxorubicin. Table 5 shows the binding energies of the compounds. The computed binding score energy values range from – 6.4244 kcal/mol (hydroxybenzoic acid) to – 14.0088 kcal/mol (doxorubicin).

Isochlorogenic acid, orientin, kaempferol, and taxifolin are the top-scoring compounds with pose scoring values of – 13.200103, – 13.5655, – 11.836662, and – 11.228658 kcal/mol, respectively. Isochlorogenic acid and orientin (flavonol glycoside) showed the best interactions with the receptor, with the lowest binding energy values of – 13.2001 kcal/mol (RMSD = 1.46 Å) and – 13.5655 kcal/mol (RMSD = 1.16 Å), respectively. These values are close to that of doxorubicin (– 14.0088 kcal/mol, RMSD = 1.71 Å), indicating that they could inhibit the receptor. Isochlorogenic acid interaction with the receptor involved hydrogen bonds with Thr71 (H-donor), Gln64 (H-acceptor), and Arg56 (H-acceptor) in addition to ionic interactions with Arg 56AA (Fig. 8). Orientin formed interactions with Ser73AA (H-donor) and π -H interaction with Glu 76 (Fig. 9).

Kaempferol (Fig. 10) formed hydrogen bonds with Gly75AA (H-donor), and taxifolin (Fig. 11) interacted with Gly75AA and Ser73AA (H-donor). The cytotoxicity of the tested extract may be attributed to the phenolic compounds and flavonoids with the best binding affinity in the molecular docking study.

The previous experimental studies are matched with our molecular docking results. Chlorogenic acid was reported to exhibit potential effects on cytotoxicity and inhibited human colon cancer cell proliferation through cell-cycle arrest and apoptosis ⁹⁵. In addition, orientin exhibited remarkable cytotoxicity and antiproliferative activity against HT-29 colon cancer cells, induced G0/G1 cell cycle arrest, regulated cyclin and cyclin-dependent protein kinases, and mediated apoptosis in human colorectal cancer HT-29 cells ⁹⁶. Further, taxifolin showed human colorectal cancer cell growth arrest in the G2 phase of the cell cycle and apoptosis in a concentration-dependent approach ⁹⁷. Besides, kaempferol's cytotoxic effects and induction of apoptosis in different human colorectal cancer cell lines have been reported ⁹⁸.

No	Compound	Binding score (kcal/mol)	RSMD_refine (Å)	E_conf (kcal/mol)
1	Cinnamic acid	- 6.7353	1.07	- 75.77
2	Methyl gallate	- 8.4939	1.36	- 99.08
3	Taxifolin	- 11.2287	1.32	- 17.49
4	7-Methoxy-4-methylcoumarin	- 6.5481	1.12	34.50
5	Quercetin	- 10.6524	0.92	35.93
6	epicatechin	- 10.6177	1.26	23.60
7	Catechin	- 10.3107	1.28	23.49
8	Lycopsamine	- 8.8984	1.89	9.61
9	Heliospathine	- 7.9719	0.88	- 7.98
10	Trachelanthamine	- 8.5378	1.44	18.25
11	Protocatechuic acid	- 8.0570	0.98	- 71.46
12	Stearidonic acid	- 7.8434	1.81	- 53.25
13	Eriodictyol	- 10.5128	1.31	9.94
14	Kaempferol	- 11.8367	0.66	- 52.05
15	Europine	- 9.2788	1.29	39.07
16	Heliosupine	- 8.5358	1.06	117.43
17	Retronecine	- 7.7943	1.82	- 60.09
18	Maleic acid	- 7.1007	0.66	- 177.99
19	Caffeic acid	- 9.5269	1.34	- 96.22
20	2-Ethoxy-4,5-dihydroxybenzaldehyde	- 9.6644	0.65	16.05
21	6-Deoxycochinolide	- 8.2384	0.89	- 1.10
22	2-Isopropylmalic acid	- 9.5676	0.84	- 36.78
23	7-Hydroxy-1-methylenepyrrolizidine	- 7.5278	0.90	4.06
24	4-Hydroxybenzoic acid	- 6.4245	1.48	- 68.26
25	Succinic acid	- 6.8769	0.83	- 176.13
26	4-Methoxycinnamic acid	- 7.3200	1.52	- 72.06
27	Filifolinoic acid	- 8.7365	0.84	3.57
28	Umbelliferone	- 9.2712	1.42	12.29
29	4,5-di-O-caffeoylquinic acid (Isochlorogenic Acid)	- 13.2001	1.46	- 4.03
30	3-O-p-coumarylquinic acid	- 9.5033	1.09	10.87
31	3-O-Feruloylquinic acid	- 10.1228	0.85	23.34
32	Orientin	- 13.5655	1.16	53.38
33	Tournefolin C	- 9.0987	0.97	9.53
34	Doxorubicin	- 14.0088	1.71	- 79.03

Table 5. Docking scores of identified compounds against colon cancer antigen 10 (PDB ID: 2HQ6).

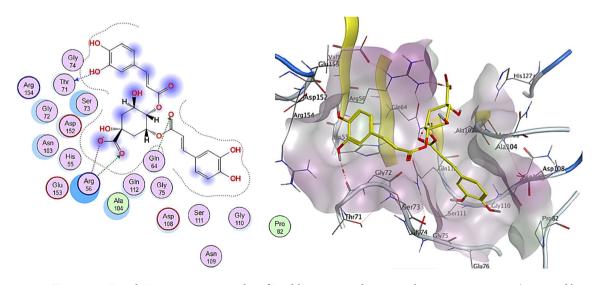


Figure 8. 2D and 3D interactions complex of Isochlorogenic acid against colon cancer antigen 10 (generated by using Molecular Operating Environment, MOE, 2014.0901⁹⁹).

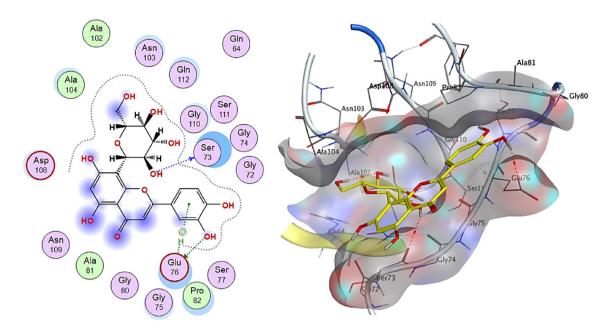


Figure 9. 2D and 3D interactions complex of Orientin against colon cancer antigen 10 (generated by using Molecular Operating Environment, MOE, 2014.0901⁹⁹).

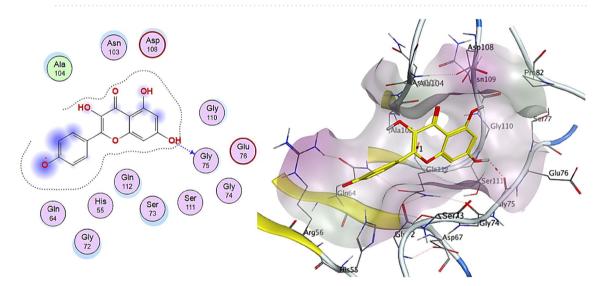


Figure 10. 2D and 3D interactions complex of Kaempferol against colon cancer antigen 10 (generated by using Molecular Operating Environment, MOE, 2014.0901⁹⁹).

Conclusion

In conclusion, the phytochemical screening of the total methanolic extract of the aerial parts of *H. ramosis-simum* (Lehm.) DC. revealed the presence of alkaloids, terpenoids, saponins, tannins, flavonoids, coumarins, polyphenolics, and reducing sugars. It is worthy to mention that all the identified compounds in the *n*-hexane fraction are the first to be reported in the genus *Heliotropium*, in addition, thirty-two compounds of different chemical classes were tentatively identified by LC–ESI–MS/MS analysis. Besides, our findings revealed that the total methanolic extract of *H. ramosissimum* aerial parts exhibits a potent in vitro antitumor effect against several cell lines especially colorectal carcinoma (Colo-205) in a dose-dependent manner and with selectivity in comparison to normal cell lines (OEC). The methanolic extract reduces the viability of invasive Colo-205 cell lines by inducing apoptosis and/or necrotic cell death. Furthermore, a docking study for the identified compounds from the LC–ESI–MS/MS analysis to find out the candidates responsible for the cytotoxic activity where isochlorogenic acid and orientin showed the highest binding scores with colon cancer antigen 10. Our findings can aid in the creation of a new alternative candidate for the selective treatment of early stages of colorectal cancer with safety on normal cells. Future studies are needed to isolate the active constituent/constituents responsible for this potent cytotoxic activity.

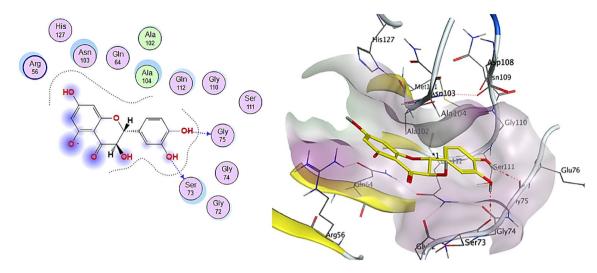


Figure 11. 2D and 3D interactions complex of Taxifolin against colon cancer antigen 10 (generated by using Molecular Operating Environment, MOE, 2014.0901⁹⁹).

Data availability

All data generated or analyzed during this study are included in this published article [and its supplementary information files].

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