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Evaluation of the anticancer effects exerted by 5-fluorouracil and heme oxygenase-1 inhibitor hybrids in HTC116 colorectal cancer cells

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ABSTRACT

Colon cancer remains a clinical challenge in industrialised countries. Its treatment with 5-Flurouracil (5-FU) develops many side effects and resistance. Thus, several strategies have been undertaken so far, including the use of drug cocktails and polypharmacology. Heme oxygenase-1 (HO-1) is an emerging molecular target in the treatment of various cancers. We recently demonstrated that a combination of HO-1 inhibitors with 5-FU and the corresponding hybrids SI1/17, SI1/20, and SI1/22, possessed anticancer activity against prostate and lung cancer cells. In this work, we evaluated these hybrids in a model of colon cancer and found that SI1/22 and the respective combo have greater potency than 5-FU. Particularly, compounds inhibit HO-1 activity in cell lysates, increase ROS and the expression of HO-1, SOD, and Nrf2. Moreover, we observed a decrease of pro-caspase and an increase in cleaved PARP-1 and p62, suggesting apoptotic and autophagic cell death and potential application of these drugs as anticancer agents.

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KEYWORDS

Hybrids; mutual prodrugs; 5-fluorouracil; heme oxygenase 1; HTC116 colorectal cancer cells

Introduction

A novel target for anticancer therapy is the HO family of enzymes (HO-1, HO-2, and HO-3) devoted to the catabolism of heme¹, and the production of iron, carbon monoxide, biliverdin, and bilirubin^{2,3}. Although HO has mainly a cytoprotective role, overexpression of the inducible isoform HO-1 has been observed in many types of cancers, contributing to cancer spread and invasiveness⁴; therefore, HO-1 inhibitors have been developed in the last years as new anticancer agents both alone and combined with chemotherapeutic drugs⁵⁻⁹.

Colorectal cancer (CRC) is the third most common cancer diagnosed, and the second cause of cancer death¹⁰. In CRC, HO-1 has been demonstrated to be involved in cancer induction and spreading. Different studies pointed out a remarkable high HO-1 expression and activity in tumour tissue from CRC patients¹¹ and, more specifically, the highest HO-1 expression has been detected in well-differentiated tumour areas of CRC. High nuclear localisation of HO-1 was also shown in human HCT116 cells where it activates transcription factors responsible for protection from oxidative stress and involved in cell proliferation, such as AP-1, AP-2, and Brn-3¹². CRC is treated with different chemotherapeutic drugs, including 5-fluorouracil (5-FU), capecitabine, irinotecan, oxaliplatin, trifluridine and tipiracil¹³. Despite the substantial advances achieved in the last years with chemotherapeutic agents against CRC, severe side effects, toxicity, and drug resistance are the major clinical problems^{14,15}.

5-FU is an essential medicine on the WHO list¹⁶, belonging to the chemical class of pyrimidine antimetabolites¹⁷; it is prescribed as first-line therapy for CRC^{18,19}. 5-FU exerts its cytotoxic activity through the inhibition of thymidylate synthase and the incorporation of its metabolites into nucleic acids, causing DNA damage; however, this action is not selective against cancer cell²⁰⁻²². In addition, unfavourable pharmacokinetic (PK) properties negatively contribute to the 5-FU drug profile²³. To overcome these limits, 5-FU is often used in combined therapy for the treatment of CRC (e.g. FOLFOX and FOLFIRI)13, but still exhibits severe cytotoxic effects and some drawbacks²⁴.

In recent years, combination therapy often used in cancer treatment is being replaced by the use of hybrids or multitarget compounds. These last are novel molecules that, combining in a single chemical entity two or more chemical features needed to act on different molecular targets, work as combined therapy reducing limitations related to combination therapy²⁵. A particular type of multitarget drug is the mutual prodrug, which is designed to release the original parent drugs after in vivo administration²⁶. This approach was successfully applied in the case of 5-FU-based polypharmacology²⁷.

We recently synthesised three innovative 5-FU/HO-1 hybrids named SI1/17, SI1/20, and SI1/22 (Figure 1). The rationale for their

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SI1/09







Figure 1. Chemical structures and HO inhibitory activity in spleen and brain microsomal fractions of tested compounds.

design was to link, in the same chemical entity, the antimetabolite 5-FU and our most potent arylethanolimidazoles (i.e. SI1/09, LS4/28, and LS6/42, respectively, Figure 1) selected from an in-house library of non-competitive HO-1 inhibitors since the alcohol functional group can be easily esterified with a cleavable linker of succinic acid which in turn is linked to 5-FU. Accordingly, hybrids were developed to release the parent components after hydrolytic or enzymatic cleavage, thus acting as mutual prodrugs. These compounds, previously tested against prostate (DU145) and lung (A549) cancer cells, gave interesting results and were shown to release, in vitro, 5-FU, and HO-1 inhibitors^{28,29}.

Being 5-FU the first-choice drug for CRC and being HO-1 involved in the proliferation of CRC cells³⁰⁻³², in this work we aimed to evaluate the antiproliferative effects of the hybrids, as well as the physical combinations of 5-FU and corresponding HO-1 inhibitors (combo treatment), in a colon cancer model. Previously, we demonstrated that in colon cancer cells the targeting of intracellular redox balance commits to cell damage and death^{33,34}. Thus, some important antioxidant and apoptotic pathways were investigated to preliminarily analyse the mechanism of action of both the hybrids and combo treatments.

Materials and methods

Drugs preparation

5-FU was purchased from Fluorochem (product code F003241). Imidazole-based HO-1 inhibitors (SI1/09, LS4/28, and LS6/42)^{35,36} and key intermediates 4-[(5-fluoro-2,4-dioxo-1,2,3,4-te trahydropyrimidin-1-yl)methoxy]-4-oxobutanoic acid (5-FU-succ) were sourced from our compounds library. 5-FU/HO-1 hybrids (SI1/17, SI1/20, and SI1/22) were resynthesized as previously described with minor modifications^{28,29}. Drugs were dissolved in dimethyl sulfoxide (DMSO) to give a 20 mM stock solution, and stored at -20°C. Before use, all different stock solutions were diluted in RPMI culture medium to obtain the proper final concentrations.

Chemistry

Solvents and reagents were purchased from commercial vendors. Flash column chromatography was performed on Merck silica gel 60, 0.040-0.063 mm. Melting point was determined in an IA9200

Electrothermal apparatus equipped with a digital thermometer in glass capillary tubes and is uncorrected. The infra-red (IR) spectrum was recorded in KBr disc on a Perkin Elmer 1600 Series FT-IR spectrometer.¹H NMR spectra was recorded on a Varian Inova Unity 200 spectrometer, using dimethyl sulfoxide- d_6 (DMSO- d_6) solution with tetramethylsilane as an internal standard. Chemical shifts are given in δ values (ppm), while coupling constants (*J*) are given in hertz (Hz). Elemental analysis for C, H, and N was within \pm 0.4% of theoretical value and was performed on a Carlo Erba Elemental Analyser Mod. 1108 apparatus.

Details regarding the synthesis of the most representative compound SI1/22 is reported herein. Briefly, a mixture of 5-FU-succ (101 mg, 0.39 mmol), 3-dimethylamino-propyl)-ethyl-carbodiimide hydrochloride (74 mg, 0.39 mmol) and 4-(dimethylamino)pyridine (5 mg, 0.04 mmol) were added to a stirred suspension of LS6/42 (111 mg, 0.32 mmol) in 10 ml of acetonitrile anhydrous. The reaction was left to stir at room temperature overnight, and the solvent removed under vacuum. The crude was purified by silica gel column chromatography using a mixture of acetone-cyclohexane (9:1, v/v) as eluent to give 1-(1-{4-[(4-bromophenyl)methoxy]phenyl}-2-(1H-imidazol-1-yl)ethyl) 4-(5-fluoro-2,4-dioxo-1,2,3,4-tetrahydropyrimidin-1-yl)methyl butanedioate (33 mg), as a white solid: mp 143.5-144.0 °C; IR (KBr, selected lines): cm-1 3448, 3081, 2960, 1756, 1611, 1513, 1458, 1408, 1357, 1245, 1133, 1001, 832, 754; ¹H NMR (200 MHz, DMSO- d_6) δ 8.11 (d, J_{H-F} = 6.6 Hz, 1H), 7.60 (d, J=8.2Hz, 2H), 7.51 (s, 1H), 7.41 (d, J=8.0Hz, 2H), 7.26 (d, J=8.3Hz, 2H), 7.11 (s, 1H), 6.99 (d, J=8.3 Hz, 2H), 6.84 (s, 1H), 5.90 (t, J=6.1 Hz, 1H), 5.55 (s, 2H), 5.09 (s, 2H), 4.34 (d, J=7.0 Hz, 2H), 2.58 (s, 4H). Anal. Calcd. for C₂₇H₂₄BrFN₄O₇: C, 52.70; H, 3.93; N, 9.10. Found: C, 52.53; H, 3.92; N, 9.08.

Measurement of HO-1 enzymatic activity

HO-1's enzymatic activity was measured in cell lysates as the difference in absorbance between 464 and 530 nm of the bilirubin produced. Cells were cultured in RPMI supplemented with 10% FBS and 1% pen-strep solution, then cells were treated for 48/72 h with compounds (LS6/42 - SI1/22) at a concentration of 5 µM. After treatment cells were harvested and HO-1 enzymatic activity was assessed. The reaction mixtures consisted of 20 mM Tris-HCl at pH 7.4 (2mg/mL), the cell lysate, 0.5-2mg/mL biliverdin reductase, 1 mM NADPH, 2 mM glucose 6-phosphate (G6P), 1 U G6P dehydrogenase and 25 µM hemin. Incubation was carried out in a circulating water bath in the dark for 1h at 37°C. The reaction was stopped by adding chloroform. After recovering the chloroform phase, the amount of bilirubin that was formed was measured with a double-beam spectrophotometer at OD 464-530 nm (extinction coefficient: 40 mM/cm⁻¹ for bilirubin). One unit of the enzyme was defined as the amount of enzyme catalysing the formation of 1 nmol of bilirubin/mg protein/h.

Cells and culture conditions

HCT116 colon cancer cells were obtained from Interlab Cell Line Collection (ICLC, Genoa, Italy) (accession number ICLC HTL95025). Cells were maintained in RPMI1640 medium enriched with 2 mM glutamine and 10% of heat-inactivated foetal bovine serum (FBS). A solution of 100 U/mL penicillin and 100 μ g/mL streptomycin was added to the medium. Cells were incubated at 37°C, in the presence of 5% CO₂ and humidified atmosphere. Once cell confluence was reached, cells were seeded in opportune culture plates at the

density of 2×10^4 cells/cm² and exposed to the different drugs or vehicle (DMSO) alone (indicated in the text as control) for the established time. DMSO concentration never exceeded 0.3% (v/v), a concentration that did not affect HCT116 cells viability.

Cell viability assay

The colorimetric assay using MTT (3–(4,5-dimethylthiazol-2-yl)-2,5diphenyltetrazolium bromide (Merck-Sigma Aldrich, MI, Italy), was employed to measure cellular metabolic activity as an indicator of cell viability and proliferation, as previously described³⁷. For the experiments, cells were seeded in 96-well plates, treated with the compounds at different concentrations (5, 10, or 15 μ M) or vehicle alone (control) for the indicated time, and incubated with 5 mg/ml of MTT for 2 h. The formazan crystals were dissolved in a lysis solution and the staining intensity was quantified by measuring the absorbance at 570 nm at a microplate reader (OPSYS MR, Dynex Technologies, Chantilly, VA, USA).

Cell cycle analysis

To analyse cell cycle distribution, after treatment with the compounds, cells were harvested by trypsinization (0.025% trypsin-EDTA; Life Technologies Ltd, Monza, Italy) and resuspended in a hypotonic solution containing $25\,\mu$ g/ml propidium iodide, 0.1% sodium citrate, 0.01% Nonidet P-40 and $10\,\mu$ g/ml RNase A for 2 h at 4° C. Cell cycle distribution was evaluated by using a FACScanto cytometer and data elaborated by Flowing software.

Western blotting analysis

After treatment, cells were lysed in RIPA buffer and protein extracted were quantified using Bradford method. Equal amount of proteins (30 µg) was separated by SDS-PAGE, electrotransferred to a nitrocellulose membrane and exposed to specific primary antibodies. HO-1 (GTX101147) antibody was purchased from Gene Tex (GeneTex, Prodotti Gianni, MI, Italy); anti-SOD (sc-133); anti-p62 (P0068) and anti-Actin (A4700) from Merck Millipore (Merck-Sigma Aldrich, MI, Italy); anti Caspase-3 (96625) and anti Nrf-2 (sc-518033) from Cell Signalling Technology (Danvers, MA, USA); anti-PARP1 from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Molecular weight markers were from Biogenerica, Catania, Italy (catalogue number RPN 800E rainbow molecular weight markers, full-range Mr 12000-225000). After incubation with specific HRP-conjugated secondary antibodies (Merck-Sigma Aldrich, MI, Italy), the blots were developed using the ECL chemiluminescent labelling systems. Optical densities of the bands were analysed by using ImageJ Software. In some cases, proteins were analysed on the same filter after stripping and re-hybridization with the specific antibodies.

ROS evaluation

The reactive oxygen species (ROS) produced after treatment were detected through the oxidation of the cell-permeant 2',7'-dichlorodihydrofluorescein diacetate (H₂DCFDA) (Molecular Probe, Life Technologies, Eugene, OR, USA) dye, as reported before³⁸. Cells were seeded in 24-well plates at the density of 2×10^4 /cm², treated and stained with 2 μ M H₂DCFDA dye for 15 min in the dark and in the presence of 5% CO₂ at 37° C. After washing, the fluorescent 2',7'-dichlorofluorescein (DCF), produced by intracellular oxidation, were analysed under a fluorescence microscope using FITC filter. Images were acquired by using an Optika IM3F fluorescence microscope equipped with camera.

Statistical analysis

Data, expressed as mean \pm SD, were evaluated with Student's t-test using Microsoft Excel software. The analysis of multiple groups of samples was performed with ANOVA test. *P* values <0.05 was considered statistically significant.

Results and discussion

Cytotoxicity and morphology of HCT116 CRC cells after treatment with 5-FU, HO-1 inhibitors, 5-FU/HO-1 hybrids and respective combo

Initially, we evaluated the effect of HO-1 inhibitors and the respective 5-FU/HO-1 hybrids in HCT116 cancer cells, which represent a human colon cancer model widely employed in therapeutic research and drug screening. Compounds were tested by performing an MTT assay to evaluate their effect on HCT116 cell viability. Table 1 shows a screening over time of the effects exerted by the compounds employed at 5 μ M concentration. The tested dose was chosen according to previous findings on the significant antiproliferative activity against different cancer cells exerted by HO-1 inhibitors at a similar concentration^{8,29}. The HO-1 inhibitor LS6/42 and the corresponding SI1/22 hybrid resulted in the most

Table 1. Time course of HCT116 cell viability after treatment with $5\,\mu\text{M}$ of the different drugs at different times.

	% of Cell viability±SD			
Compound	24 h	48 h	72 h	
Control	100±2.2	100 ± 4.3	100±2	
LS6/42	78±7.2	70 ± 5.5	67±6.1	
SI1/22	85 ± 4.7	80 ± 3.5	77 ± 5.5	
SI1/09	110 ± 6.6	100 ± 5.7	101 ± 4.5	
SI1/17	98 ± 7.5	90±3.7	85 ± 5.7	
LS4/28	98±8.2	95±5.1	92±5.4	
SI1/20	100 ± 9.5	100 ± 4.3	93±8.5	

MTT assay was carried out as reported in the methods. Viability was calculated as the percentage of control (HCT116 cells maintained in the presence of vehicle alone) \pm SD.

efficacious molecules in inhibiting HCT116 cell viability, after 72h of treatment.

The analysis of cell cycle distribution profiles of more efficacious compounds showed that SI1/22 induced an increase in the pre-G0/G1 population seen to the left of the G0/G1 peak, indicating DNA fragmentation that can be associated with apoptotic cell death. Differently, LS6/42 treatment induced an increase of G2/M peak which was accompanied by a moderate increase of pre-G0/ G1 peak (Figure 2).

These promising data prompted us to investigate the action of novel compounds in more detail and the effects of the treatment using HO-1 inhibitors alone or in combination with 5-FU on HCT116 cell viability (Figure 3). As a general trend, HO-1 inhibitors showed a low to moderate effect on cell viability, while the hybrids produced a significant dose-dependent effect comparable to that of 5-FU. Among the tested hybrids SI1/22, bearing a bulkier and more hydrophobic 4-bromobenzyloxy group at the para-position of the phenylethyl moiety, exerted better antiproliferative activity than the 3-bromo (SI1/17) and the 3-phenyl (SI1/20) analogs, respectively; also compared to the reference drug 5-FU. As a result, a correlation between increased lipophilicity and higher antiproliferative potency was observed, likely due to the promotion of cellular uptake and considering the higher predicted permeability value of SI1/22 (Table 2). Particularly, SI1/22 was more potent than 5-FU at all tested concentrations resulting in the best compound among the series, as well as the combo administration of LS6/42 and 5-FU resulted in even more effective in terms of antiproliferative action (Figure 3A). Also, agreeing with extensive structure-activity relationship studies (SARs) performed on different azole-based analogs, and agreeing with the solved crystal structure of the protein in complex with non-competitive HO-1 inhibitors (e.g. QC-80)³⁹, the simultaneous accommodation of bulky groups (i.e. 5-FU-succinyl and 4-bromobenzyloxy moiety as for SI1/22) into the two main hydrophobic pockets of HO-1 is generally well-tolerated and seems to be an optimal combination for stabilising the inactive conformation of the enzyme.

Based on these results, the HO-1 inhibitor LS6/42 alone or in combination with 5-FU and the corresponding hybrid SI1/22 were selected for subsequent experiments. The morphological observation of the treated cells with 15 μ M of each compound, and the corresponding drug combination, after 72 h exposure, showed that LS6/42 did not induce a significant morphological change in cancer cells, while SI1/22 and the LS6/42+5-FU combo produced a significant reduction of cell density and a slight morphological



Figure 2. Flow cytometry analysis of the cell cycle phase distribution of HCT116 cells after 72 h incubation with 5 µM SI1/22 or LS6/42. Hypotonic propidium iodide staining was used to evaluate DNA content as described in Methods employing Flowing software.



Figure 3. Effect of 5-FU, HO-1 inhibitors, hybrids, and combo treatment on HCT116 cell viability and morphology. (A) Evaluation of cell viability by MTT assay after 72h of treatment using different concentrations of the drugs. The results represent the mean \pm SD of almost three separate experiments done in triplicate. Significant *vs* untreated control cells: *p < 0.05; **p < 0.01; ***p < 0.001. (B) Representative images of HCT116 cells after 72h of treatment with 15 μ M of the compounds. Cells were visualised under a light microscope (200× magnification) and the pictures were acquired by IM50 Leica Software (Leika Microsystems, Wetzlar, Germany).

 Table 2. Predicted lipophilicity/permeability and experimental antiproliferative effects in HCT116 cells of hybrids.

Compound	Lipophilicity (SK logD)	Permeability (Papp, nm/s)	% of cell death, 72h [5μM]
SI1/22	2.97	24.4	23
SI1/17	1.45	20.6	15
SI1/20	2.28	21.8	7

Selected ADME properties were predicted using PreADMET web-based application (http://preadmet.bmdrc.kr).

variation, as the cells appeared more rounded and less close to one another, thus supporting an anti-proliferative effect (Figure 3B).

Effect on HO-1 enzymatic activity, ROS production and Nrf2dependent antioxidant pathways

Then, we wanted to confirm that hybrid SI1/22 was able to inhibit HO-1 activity not only in microsomal preparation but also in cell lysate, an experimental condition that better reproduces a cell

Table 3. Inhibition of HO-1 enzymatic activity in HCT116 treated cells with LS6/42 and SI1/22.

	picomol/mg prot/1 h	% inhibition	
Compound	[5 µM], 72 h		
Control	53.5 + 1.118	-	
_S6/42	11.42+0.903	(78.65%)	
511/22	17.67 + 1.121	(66.9%)	

Data are representative of at least three independent experiments,

environment. For this purpose, bilirubin formation was measured in HCT116 cell lysates after 72h of treatment. SI1/22 was able to reduce HO-1 activity similarly to the positive control LS6/42 (i.e., 78.65% vs. 66.9%) (Table 3). This result suggests that hybrid SI1/22 can cross cellular membranes and then it is cleaved to regenerate parental drugs and inhibit HO activity.

Additionally, to correlate HO-1 inhibition to oxidative stress conditions, ROS levels and antioxidant proteins expression were evaluated. Results depicted in Figure 4 clearly show that, after 3 h exposure, ROS levels highly increased after both treatments with hybrid SI1/22 and the LS6/42+5-FU combo, while no remarkable



Figure 4. Effect of 5-FU, HO-1 inhibitors, hybrids, and combo treatment on oxidative stress. Assessment of ROS levels in HCT116 cells treated with 15 µM of tested drugs for 3 h; the images were acquired under a fluorescence microscope at 200 x magnification.

differences were observed for LS6/42 and 5-FU as a stand-alone treatment. Although 5-FU is known to increase ROS cellular levels⁴⁰, there was no increase in our experimental conditions. On the other hand, hybrid SI1/22 and combo treatment were able to significantly induce ROS production even at low concentrations and after just 3 h of exposure.

Consequently, due to the high and early increase in ROS levels, the activation of an antioxidant response was analysed by measuring the expression levels of two antioxidant enzymes involved in oxidative stress, such as HO-1 and SOD. Western blotting and densitometric analysis highlighted an increase in both enzyme expressions compared to the control group (Figure 5A). These data were consistent with the previously observed ROS production with a major effect detected in the case of LS6/42+5-FU combo treatment.

Therefore, the modulation of antioxidant pathways was analysed by measuring the expression level of Nrf2, which is one of the major transcription factors that promotes cellular defense against oxidative stress through the expression of many antioxidant genes including *HMOX1* encoding for HO-1⁴¹. Collected data depicted in Figure 5B showed an increased Nrf2 level when cells were treated with both hybrid Sl1/22 and 5-FU+LS6/42 combo compared to the control, in accordance with the increase of HO-1 levels (Figure 5A).

Involvement of apoptotic or autophagic mechanism in drugsinduced cell death

To investigate what type of cell death pathway was induced by the compounds, we first analysed the levels of pro-caspase 3 as indicative of its activation to caspase-3. Caspase-3 is a well-known protease that catalyses the specific cleavage of many key proteins responsible for apoptosis. Western blotting and densitometric analysis suggested that all treatments decreased the pro-caspase-3 expression level in HCT116 cells, suggesting a possible promotion of apoptosis through the caspase-3-dependent pathway (Figure 5A). Active caspase-3 is known to cleave PARP1, a nuclear protein involved in DNA damage repair systems⁴². PARP1 and cleaved-PARP1 levels were also evaluated, showing a significant increase in cleaved isoform especially after SI1/22 and combo treatments (Figure 5C).

Additionally, to further investigate the mechanism of action of the hybrid compound we analysed p62 expression, a protein involved in ROS-dependent cell stress and autophagic process⁴¹.

Interestingly, treatment with LS6/42, SI1/22, and LS6/42+5-FU combo highly induced p62 accumulation in HCT116 cells (Figure 5A), suggesting an involvement of autophagy in the mechanism of action of the compounds. Since the autophagic pathway can play an opposite role (pro- or anti-survival) in cells⁴³, it will be of interest to investigate deeper how the hybrids or the combination influence autophagy in colon cancer cell models.

Conclusions

In this work we aimed to evaluate whether the 5-FU/HO-1 hybrids SI1/17, SI1/20, and SI1/22 previously developed as antiproliferative agents against prostate and lung cancer cells, might work also against CRC, being this last treated with 5-FU as a first-choice drug. Here we have demonstrated that the simultaneous use of an inhibitor of HO-1 and 5-FU, both as a combo treatment or as resulting from hybrid cleavage, in the case of SI1/22, exerted antiproliferative activity greater than that 5-FU itself, thus allowing the use of lower and therefore potentially less toxic doses. Despite cells increased expression of HO-1, SOD, and Nrf-2 in response to the oxidative stress, the inhibition of HO-1 activity is maintained in the cellular environment after treatment with our compounds. Our preliminary study on the mechanism of cell death has highlighted that apoptosis and autophagy are involved, paving the way for new applications in the development of innovative CRC treatments.

Author contributions

Conceptualisation, M.G. and S.I.; methodology, L.S., A.N., V.S., M.G., S.I.; validation M.G. and S.I.; synthesis, purification, and compounds characterisation, L.S., V.P., S.I.; contributed reagents, materials, and analysis tools, V.S., L.V., M.G., S.I.; formal analysis, L.S., V.S. M.G., S.I.; investigation, A.N., V.C., F.A., V.P., L.V.; data curation, L.S., A.N., F.A., V.C., M.G., S.I.; writing original draft preparation, L.S., M.G., S.I.; writing, review and editing, L.S., M.G., S.I.; visualisation, A.N., L.V., V.C.; supervision, L.S., M.G., S.I.; funding acquisition, M.G. and S.I. All authors have read and agreed to the published version of the manuscript.

Disclosure statement

The authors declare no conflict of interest.



Figure 5. Analysis of apoptosis and autophagy-related proteins. Representative images and relative densitometric analysis of western blotting against different proteins. HO-1, SOD, pro-caspase-3, p62 (A) and PARP (C) protein expression levels were evaluated after 72 h of treatment, Nrf2 (B) protein after 48 h. The histograms are related to relative intensities over actin. *p < 0.05; **p < 0.01; ***p < 0.001 vs untreated cells (control).

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Data availability statement

The data that support this study are available from the corresponding authors upon reasonable request.

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