

# *Achillea millefolium* L. hydroethanolic extract inhibits growth of human tumor cell lines by interfering with cell cycle and inducing apoptosis

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## ABSTRACT

The cell growth inhibitory activity of the hydroethanolic extract of *Achillea millefolium* was studied in human tumor cell lines (NCI-H460 and HCT-15) and its mechanism of action was investigated. The GI<sub>50</sub> concentration was determined with the sulforhodamine B assay and cell cycle and apoptosis were analyzed by flow cytometry following incubation with PI or Annexin V FITC/PI, respectively. The expression levels of proteins involved in cell cycle and apoptosis were analyzed by Western blot. The extracts were characterized regarding their phenolic composition by LC-DAD-ESI/MS. 3,5-O-Dicaffeoylquinic acid, followed by 5-O-caffeoylquinic acid, were the main phenolic acids, while, luteolin-O-acetylhexoside and apigenin-O-acetylhexoside were the main flavonoids. This extract decreased the growth of the tested cell lines, being more potent in HCT-15 and then in NCI-H460 cells. Two different concentrations of the extract (75 and 100 µg/mL) caused alterations in cell cycle profile and increased apoptosis levels in HCT-15 and NCI-H460 cells. Moreover, the extract caused an increase in p53 and p21 expression in NCI-H460 cells (which have wt p53), and reduced XIAP levels in HCT-15 cells (with mutant p53). This work enhances the importance of *A. millefolium* as source of bioactive phenolic compounds, particularly of XIAP inhibitors.

## 1. Introduction

Cancer is one of the most lethal diseases in the world, being the leading cause of death in economically developed countries and the second one in developing countries. It is crucial to find new efficient therapies to treat this disease (Holohan et al., 2013). Plants were traditionally used in the past to treat cancer and nowadays, especially in developing countries, a large percentage of the population is still using plants instead of the pharmaceutical products available (Uniyal et al., 2006). Therefore, the presence of bioactive compounds in plants and other natural products has been extensively investigated by many scientists, in order to apply this knowledge to the treatment of cancer.

*Achillea millefolium*, also known as yarrow, belongs to the Asteraceae family and is a traditional plant that can be found in several countries of Europe, Asia, North Africa and North America (Potrich et al., 2010; Vitalini et al., 2011). This plant has been used during centuries to treat several conditions, such as gastrointestinal and hepatobiliary disorders,

inflammation and diabetes (Akram, 2013; Benedek and Kopp, 2007). Its sedative properties were also described, since recent studies have shown that a hydroalcoholic extract of *A. millefolium* has anxiolytic effects *in vivo* (Baretta et al., 2012). In addition, this plant can be used in a distillate solution as a disinfectant due to its antibacterial effect (Miranzadeh et al., 2015). Interestingly, some studies showed the antitumor potential of some extracts of *A. millefolium* in human tumor cell lines (Csupor-Löffler et al., 2009; Dias et al., 2013). In the case of aqueous and methanolic extracts, the bioactivity was related with phenolic compounds, mainly phenolic acid derivatives (caffeoylquinic acid derivatives) and flavones (Dias et al., 2013).

The aim of this work was to investigate the tumor cell growth inhibitory effect of the hydroethanolic extract of *A. millefolium* in different human tumor cell lines: NCI-H460 (non-small cell lung cancer), and HCT-15 (human colorectal adenocarcinoma). Moreover, the effects of the hydroethanolic *A. millefolium* extract on the cell cycle profile and apoptosis were also studied in these cell lines.

**Abbreviations:** FBS, Fetal Bovine Serum; PBS, Phosphate Buffer Solution; RT, Room Temperature; SRB, Sulforhodamine B

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**Table 1**  
Phenolic compounds profile of the *Achillea millefolium* L. hydroethanolic extract.

Peak	Rt (min)	$\lambda_{\text{max}}$ (nm)	Pseudomolecular ion [M-H] <sup>−</sup> (m/z)	MS <sup>2</sup> (m/z)	Identification	Quantification (mg/g extract)
1	5.2	324	353	191(100),179(67),173(5),135(43)	3-O-Caffeoylquinic acid	0.21 ± 0.01
2	6.5	324	341	179(100)	Caffeic acid hexoside	0.060 ± 0.002
3	7.3	326	353	191(54),179(86),173(100),135(33)	4-O-Caffeoylquinic acid	0.255 ± 0.003
4	8.1	328	353	191(100),179(10),173(8),135(5)	5-O-Caffeoylquinic acid	5.12 ± 0.04
5	11.4	332	593	473(21),383(10),353(25)	Apigenin-C-hexoside-C-hexoside	0.102 ± 0.003
6	15.1	334	563	473(10),443(13),383(22),353(20)	Apigenin-C-hexoside-C-pentoside	0.163 ± 0.003
7	15.4	341	563	473(12),443(21),383(12),353(24)	Apigenin-C-glucose-C-pentoside	0.101 ± 0.004
8	16.4	348	447	357(80),327(85),297(32),285(12)	Luteolin-6-C-glucoside	0.072 ± 0.001
9	17.4	354	595	301(100)	Quercetin-O-pentosyl-hexoside	0.032 ± 0.001
10	17.7	354	463	301(100)	Quercetin-O-hexoside	0.58 ± 0.03
11	18.2	344	695	651(100),609(3),447(17),301(15)	Quercetin-O-malonylhexosyl-rhamnoside	0.093 ± 0.004
12	19.5	348	579	417(8),285(42)	Kaempferol-O-pentosyl-hexoside	0.069 ± 0.002
13	19.6	352	609	301(100)	Quercetin-3-O-rutinoside	0.20 ± 0.01
14	20.5	342	593	269(100)	Apigenin-O-dihexoside	0.12 ± 0.01
15	20.6	356	477	315(100)	Isorhamnetin-O-hexoside	0.086 ± 0.002
16	21.0	328	515	353(70),335(32),191(40),179(71),173(91),161(16),135(22)	3,4-O-Dicaffeoylquinic acid	1.18 ± 0.04
17	21.4	350	505	301(100)	Quercetin-O-acetylhexoside	0.35 ± 0.01
18	22.4	352	505	301(100)	Quercetin-O-acetylhexoside	0.18 ± 0.01
19	22.6	328	515	353(96),335(4),191(100),179(71),173(8),161(12),135(20)	cis 3,5-O-Dicaffeoylquinic acid	7.55 ± 0.09
20	22.9	328	515	353(90),335(10),191(100),179(65),173(7),161(11),135(12)	trans 3,5-O-Dicaffeoylquinic acid	5.51 ± 0.01
21	23.5	340	563	269(100)	Apigenin-O-pentosyl-hexoside	0.24 ± 0.01
22	25.4	328	515	353(15),335(3),191(44),179(59),173(81),161(14),135(15)	4,5-O-Dicaffeoylquinic acid	2.20 ± 0.02
23	25.5	330	431	269(100)	Apigenin-7-O-glucoside	0.294 ± 0.002
24	26.2	348	489	285(100)	Luteolin-O-acetylhexoside	1.40 ± 0.05
25	28.3	350	519	315(100)	Isorhamnetin-O-acetylhexoside	0.039 ± 0.001
26	29.2	338	473	269(100)	Apigenin-O-acetylhexoside	0.070 ± 0.002
27	30.3	338	473	269(100)	Apigenin-O-acetylhexoside	1.14 ± 0.03
28	31.2	340	473	269(100)	Apigenin-O-acetylhexoside	0.065 ± 0.002
Total phenolic acids						22.1 ± 0.1
Total flavonoids						5.39 ± 0.01
Total phenolic compounds						27.48 ± 0.09

**Table 2**  
Determined GI<sub>50</sub> concentrations of the *A. millefolium* extracts.

Extract	GI <sub>50</sub> Concentrations (μg/mL) in different cell lines	
	NCI-H460	HCT-15
<i>A. millefolium</i> hydroethanolic extract	187.3 ± 25.7	70.8 ± 8.1

GI<sub>50</sub> corresponds to the mean ± S.E. of at least three independent experiments, each of them carried out with duplicates. Doxorubicin was used as a positive control, with the following results having been obtained: 45.8 ± 11.5 nM in NCI-H460 cells and 293.3 ± 7.2 nM in HCT-15 cells. GI<sub>50</sub> refers to the concentration that inhibits 50% of cell growth. S.E. refers to standard error.

## 2. Methods

### 2.1. Plant and preparation of the extract

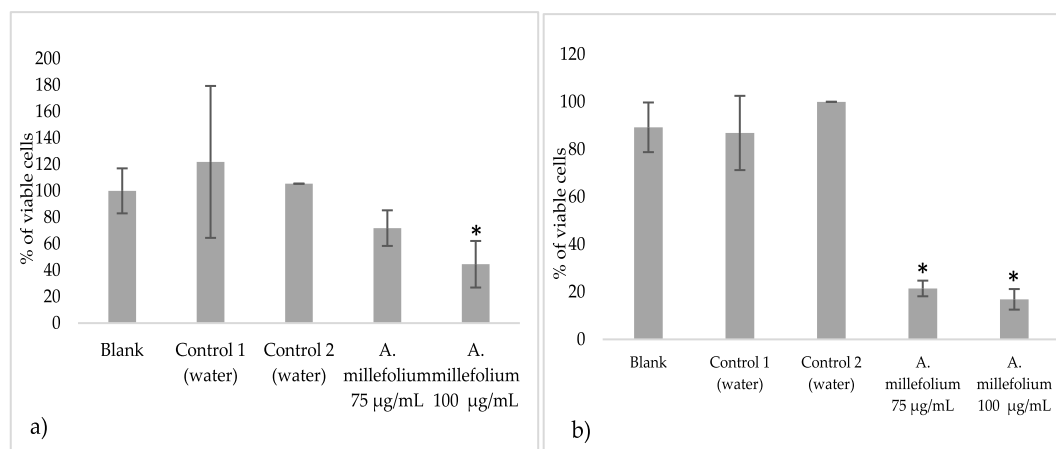
*Achillea millefolium* L. was collected in Cova de Lua, Bragança, as previously described by Dias et al. (Tozyo et al., 1994). The sample was lyophilized (FreeZone 4.5, Labconco, Kansas City, MO, USA), reduced to a fine powder (~20 mesh), homogenized and conserved in deep freezing conditions (−80 °C) for further analysis.

The hydroethanolic extract was obtained by macerating the fine and lyophilized powder (1.5 g) with two subsequent portions of ethanol/water (80:20, v/v). Each portion was centrifuged at 150 rpm for 1 h at room temperature (RT). The first portion was filtered and the solid residue was extracted again. Both filtered supernatants (Whatman No. 4 filter paper) were then evaporated under vacuum at 35 °C (Büchi R-210, Flawil, Switzerland). The remaining water residue was lyophilized (FreeZone 4.5 model 7750031, Labconco, KS, USA). The resulting fine powder was homogenized and the crude extract was stored in the dark at RT until tested. Stock solutions of the plant extract were prepared in

water and stored at −20 °C until further use.

### 2.2. Phenolic characterization of the extract

Dry lyophilized extract was dissolved in water/ethanol (80:20, v/v) in sonic bath at a final concentration of 5 mg/mL, and filtered through a 0.22 μm nylon filter for chromatographic analysis. A Dionex Ultimate 3000 UPLC (Thermo Scientific, San Jose, CA, USA) with a diode array detector (DAD, 280 and 370 nm, as preference wavelengths) and coupled to a Linear Ion Trap LTQ XL mass spectrometer (Thermo Finnigan, San Jose, CA, USA) equipped with an electrospray ionization was used to perform the analysis (Bessada et al., 2016). The identification of phenolic compounds was achieved by comparing retention times, UV–Vis and mass spectra with available standard compounds. Otherwise, available data reported in the literature were applied to tentatively identify the compounds. Quantitative analysis was performed from a 5-level calibration curve (2.5–100 μg/mL) of each available phenolic standard constructed upon the UV signal: apigenin-6-C-glucoside ( $y = 246.05x - 309.66$ ;  $R^2 = 0.9994$ ); apigenin-7-O-glucoside ( $y = 159.62x + 70.50$ ;  $R^2 = 0.999$ ); caffeic acid ( $y = 611.9x - 4.5733$ ;  $R^2 = 0.999$ ); 5-O-caffeoylquinic acid ( $y = 313.03x - 58.20$ ;  $R^2 = 0.999$ ); kaempferol-3-O-glucoside ( $y = 288.55x - 4.05$ ;  $R^2 = 1$ ); kaempferol-3-O-rutinoside ( $y = 239.16x - 10.587$ ;  $R^2 = 1$ ); luteolin-6-C-glucoside ( $y = 508.54x - 152.82$ ;  $R^2 = 0.997$ ); luteolin-7-O-glucoside ( $y = 80.829x - 21.291$ ;  $R^2 = 0.999$ ); quercetin-3-O-glucoside ( $y = 253.52x - 11.615$ ;  $R^2 = 0.999$ ) and quercetin-3-O-rutinoside ( $y = 281.98x - 0.3459$ ;  $R^2 = 1$ ). Identified phenolic compounds with unavailable commercial standard were quantified via the calibration curve of the most similar standard available. The analyses were carried out in triplicate and the results were expressed as mean values and standard deviations (SD), in mg/g of lyophilized extract.



**Fig. 1.** Effect of *A. millefolium* hydroethanolic extract on NCI-H460 (a) and HCT-15 (b) viable cell number. Cells were treated for 48 h with medium (Blank), extract (75 µg/mL and 100 µg/mL) or with the corresponding water volumes (control 1 and control 2, respectively). Viable cell number was measured with the Trypan blue exclusion assay.

### 2.3. Cell culture

Two human tumor cell lines were used in this study: NCI-H460 (non-small cell lung cancer, a kind gift from NCI, USA), and HCT-15 (human colorectal adenocarcinoma). The cells were regularly kept in RPMI-1640 medium supplemented with Ultraglutamine I and 25 mM HEPES (Lonza) and Fetal Bovine Serum (FBS, Biowest). The FBS was used at a concentration of 5% (v/v) for the Sulforhodamine B (SRB) assay and of 10% (v/v) for all the remaining assays. The cells were maintained at 37 °C in a humidified incubator with 5% CO<sub>2</sub>. The experiments described in the next sections were performed only when exponentially growing cells presented more than 95% viability determined with the Trypan blue exclusion assay. All cell lines were genotyped and regularly monitored for mycoplasma contamination by PCR.

### 2.4. Cell growth inhibition assay

In order to determine the *in vitro* cytotoxicity of the *A. millefolium* extract, the SRB (sulforhodamine B) assay was carried out in the three human tumor cell lines mentioned above.

Cells were plated in 96-well plates at an optimal density previously determined to allow exponential cell growth during the assay ( $5.0 \times 10^4$  cells/mL for H460 cells and  $1.0 \times 10^5$  cells/mL for HCT-15 cells) (Santos et al., 2013). Cells were treated 24 h later with five serial dilutions of the extract, ranging from 400 µg/mL to 25 µg/mL. As positive control, doxorubicin was used in concentrations ranging from 150 nM to 9.37 nM. As negative control, water was used (in a volume corresponding to the one used when testing the highest concentration of the extract). Two plates were prepared: one to be analyzed immediately after extract addition (T0 plate) and another one to be analyzed 48 h after extract addition (T48 plate). After 48 h of incubation for the T48 plate (or immediately for the T0 plate), cells were fixed by adding ice-cold 10% (w/v) trichloroacetic acid (TCA) (Panreac, Barcelona, Spain) and stained with 1% SRB (Sigma Aldrich, St. Louis, MO, USA) in 1% (v/v) of acetic acid (Sigma Aldrich, St. Louis, MO, USA). The bound dye was solubilized by adding 10 mM Tris base solution (Sigma Aldrich, St. Louis, MO, USA) and the absorbance was measured at 510 nm in a microplate reader (BioTek® Synergy MX, Winooski, VT, USA). The GI<sub>50</sub> concentration of the *A. millefolium* extract was determined for each cell line tested (Vichai and Kirtikara, 2006).

### 2.5. Preparation of cells for further analysis

In order to analyze the effect of the extract on cell cycle profile,

apoptosis, and on the expression of selected proteins, the NCI-H460 and HCT-15 were plated at a concentration of  $1.0 \times 10^5$  cells/well and  $2.0 \times 10^5$  cells/well, respectively, in 6-well plates. After 24 h, cells were treated with the *A. millefolium* extract at two different concentrations, 75 µg/mL and 100 µg/mL. Medium and water were added to Blank and control cells, respectively, at the same volumes as the ones used when testing the 2 extract concentrations (control 1 corresponding to the 75 µg/mL extract; control 2 corresponding to the 100 µg/mL extract). After 48 h treatment, cells were trypsinized and centrifuged at 1200 rpm for 5 min. The percentage of viable cells was determined with the trypan blue exclusion assay.

#### 2.5.1. Analysis of cell cycle profile

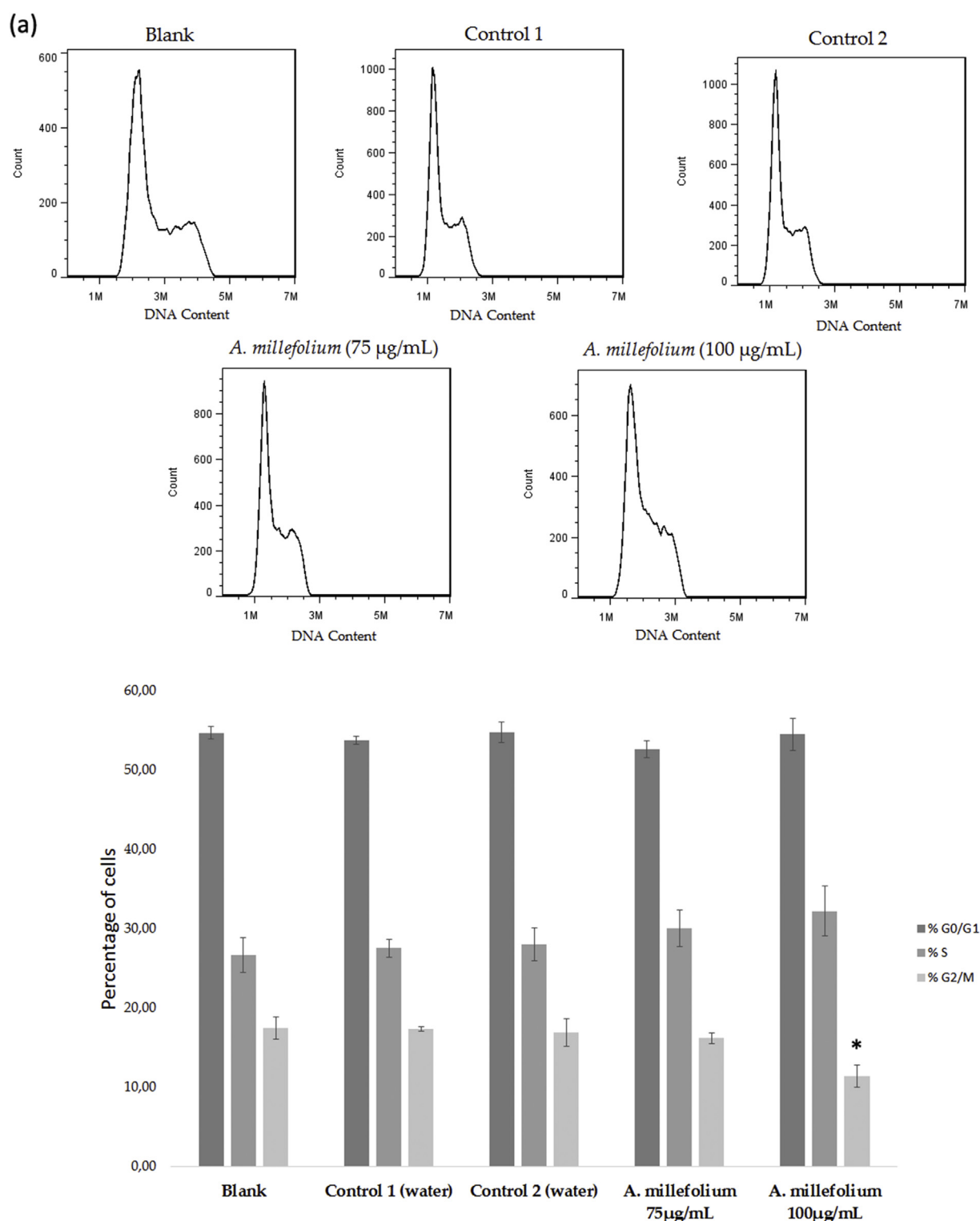
Cells were washed with a Phosphate Buffer Solution (PBS) (Sigma Aldrich, St. Louis, MO, USA), fixed with ice-cold 70% ethanol (Sigma Aldrich, St. Louis, MO, USA) and stored at 4 °C for at least 12 h. Cells were then centrifuged for 5 min at  $290 \times g$  and pellets were re-suspended in a solution of PBS containing 5 µg/mL propidium iodide (Sigma Aldrich, St. Louis, MO, USA) and 0.1 mg/mL RNase A (Sigma Aldrich, St. Louis, MO, USA). Cellular DNA content analysis was performed by flow cytometry (BD Accuri™ C6 Flow cytometer, USA) and the percentage of cells in the G0/G1, S and G2/M phases of the cell cycle was determined following cell debris and aggregates exclusion and plotting at least 10,000 events per sample, as previously described (Vasconcelos et al., 2000). All the results were analyzed using the FlowJo software (version 7.6.5, Tree Star, Inc., Ashland, USA).

#### 2.5.2. Analysis of apoptosis

Cell pellets were washed and re-suspended in PBS. After centrifugation at  $290 \times g$ , cells were analyzed for apoptosis levels with the Annexin V-FITC Apoptosis Detection Kit (eBioscience), as previously described (Vaz et al., 2012). All the samples were then analyzed by flow cytometry (BD Accuri™ C6 Flow cytometer, USA) plotting at least 10,000 events per sample. Results were analyzed using the FlowJo software (version 7.6.5, Tree Star, Inc., Ashland, USA).

#### 2.5.3. Protein expression analysis

Cellular pellets were washed in PBS and Winman's buffer (1% NP-40, 0.1 M Tris-HCl pH 8.0, 0.15 M NaCl and 5 mM EDTA) supplemented with protease inhibitor cocktail (Roche, Indianapolis, USA) was used to lyse the cells. The total protein content of lysates was quantified with a modified Lowry protocol using the DC™ Protein Assay kit (Bio-Rad, Hercules, CA, USA), following manufacturer's instructions. Protein lysates were then loaded on a 12% SDS-PAGE gel and transferred into a nitrocellulose membrane (GE Healthcare Life science, Cleveland, OH,



**Fig. 2.** Cell cycle distribution of NCI-H460 (a) and HCT-15 (b) cells treated with the hydroethanolic extract of *A. millefolium*. Cells were treated with *A. millefolium* (75 µg/mL or 100 µg/mL) during 48 h. Water (in the same volumes as the used for each treatment) was included as controls. Upper panels correspond to the histograms with cell cycle profile, following the exclusion of cellular aggregates and debris. Lower panels correspond to the % of cells in different cell cycle phases. Results are the mean  $\pm$  SE of 3 independent experiments. \* $p \leq 0.05$  between each treatment and the respective control.

USA). Membranes were then blocked in TBS-T [Tris-buffered saline solution with 0.1% Tween 20 (Promega, Fitchburg, USA)], containing 5% (w/v) non-fat dry milk (Nestlé, São Paulo, Brazil) for at least 30 min. The following primary antibodies were used: rabbit anti-PARP-1 (1:2000, sc-7150, Santa Cruz Biotechnology, Heidelberg, Germany), mouse anti-p53 (1:200, sc-126, Santa Cruz Biotechnology), mouse anti-

caspase 3 (1:1000, sc-7272, Santa Cruz Biotechnology); mouse anti-p21 (1:250, Ab-1, Calbiochem, USA), mouse anti-XIAP (1:1000, 2042, Cell Signaling Technology, Danvers, MA, USA), mouse anti-tubulin (1:10000, Sigma Aldrich, St. Louis, MO, USA) and goat anti-actin (1:2000, sc-1616, Santa Cruz Biotechnology). The secondary antibodies used were: anti-rabbit IgG-HRP, anti-mouse IgG-HRP and anti-goat IgG-

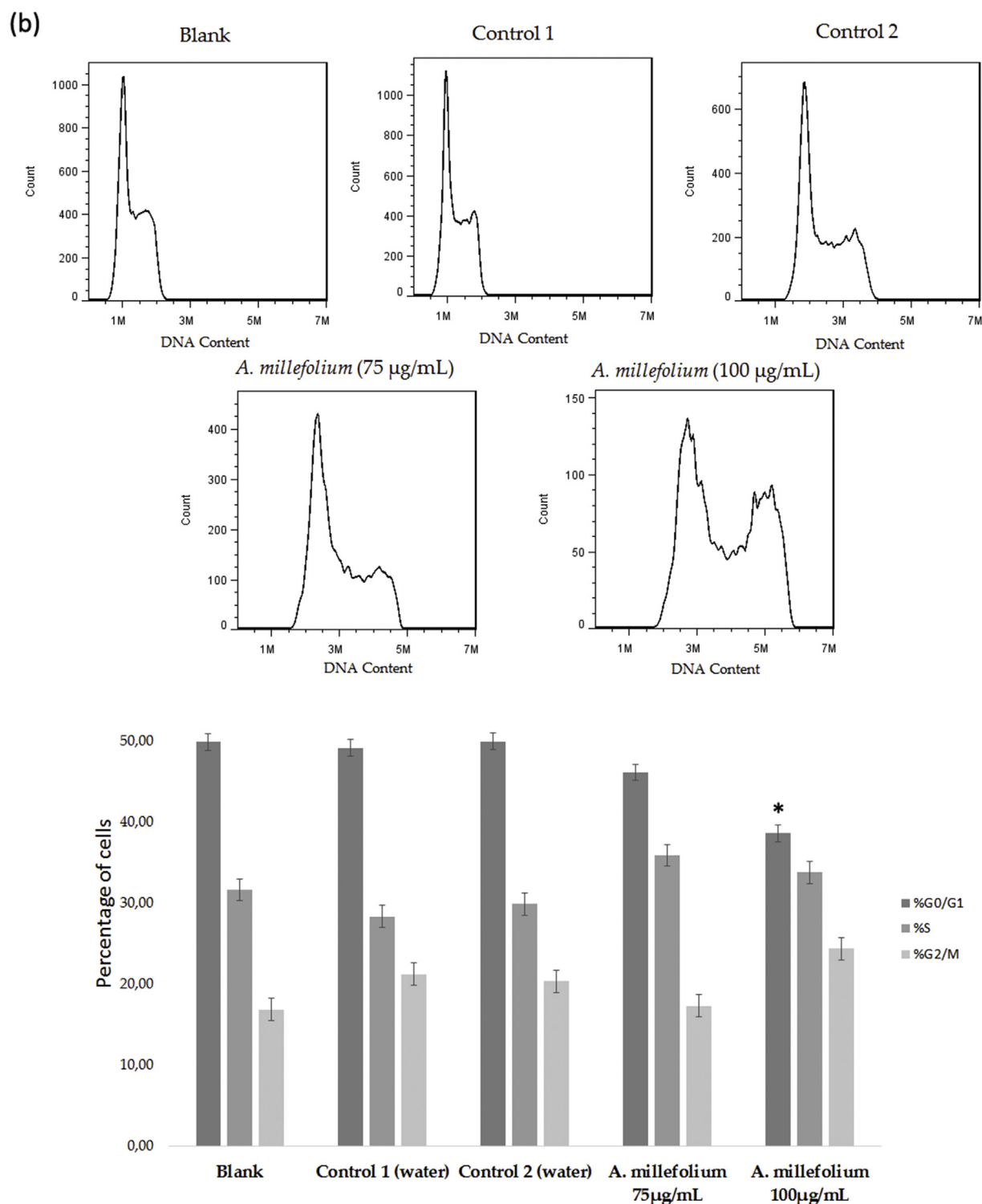


Fig. 2. (continued)

HRP (1:2000, Santa Cruz Biotechnology). The Amersham™ ECL Western Blotting Detection Reagents (GE Healthcare, Wisconsin, USA), the High Performance Chemiluminescence Film (GE Healthcare, Wisconsin, USA), and the Kodak GBX developer and fixer (Sigma Aldrich, St. Louis, MO, USA) were used for signal detection (Lopes-Rodrigues et al., 2016).

## 2.6. Statistical analysis

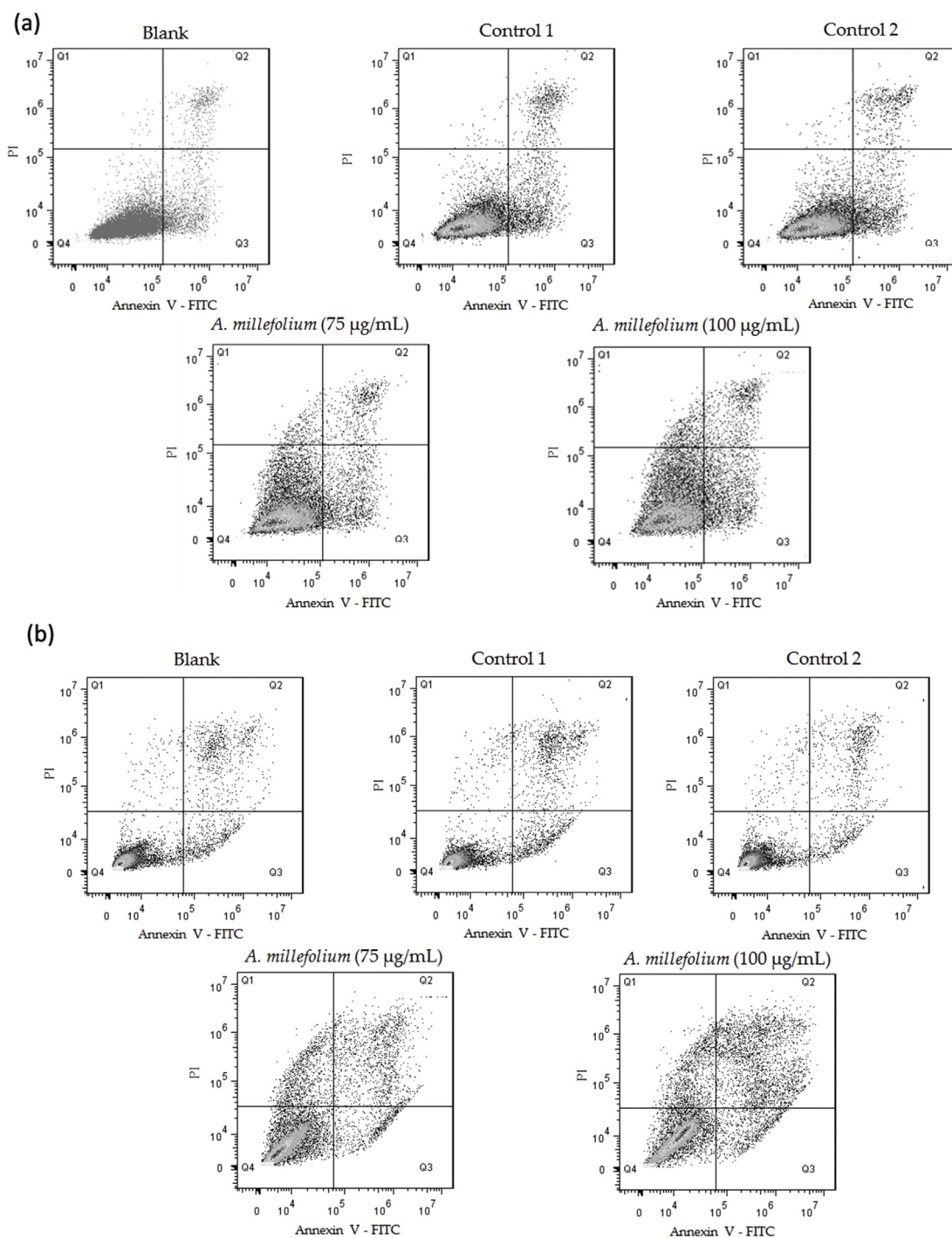
The results of at least three independent experiments were statistically analyzed using the two-tailed paired Student's t-test.

## 3. Results

### 3.1. Phenolic compounds in *A. millefolium* hydroethanolic extract

Table 1 presents the chromatographic characteristics of the phenolic compounds found in *A. millefolium* hydroethanolic extract. Twenty-eight phenolic compounds were detected, being identified 8 phenolic acid derivatives (compounds 1, 2, 3, 4, 16, 19, 20 and 22) and 20 flavonoid glycoside derivatives (compounds 5–15, 17, 18, 21, 23–28). Phenolic acids were the main class of polyphenols present, consisting of



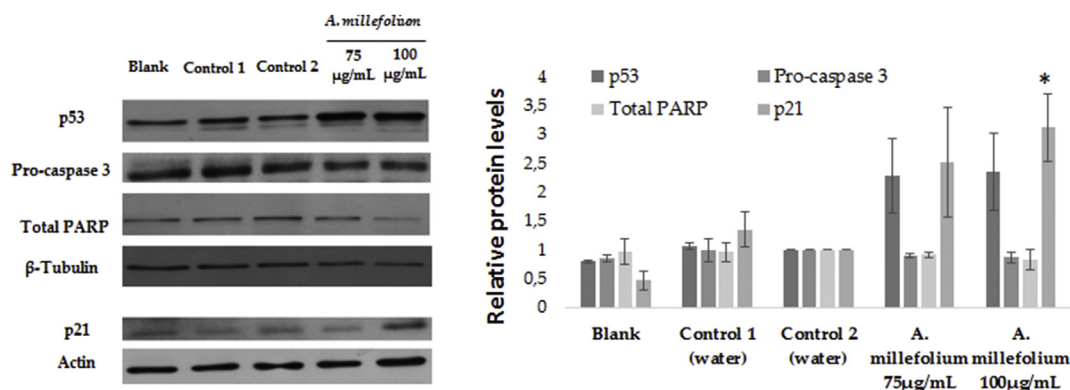


(c)

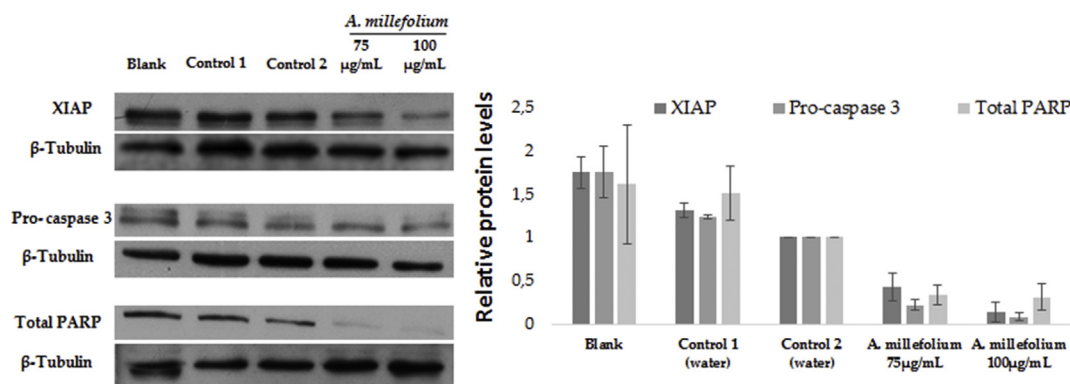
Samples	% of cells in apoptosis	
	NCI-H460	HCT-15
Blank	13.25 ± 1.53	9.82
Control 1	11.26 ± 1.67	9.21 ± 0.74
Control 2	9.81 ± 1.18	8.96 ± 1.69
<i>A. millefolium</i> (75 µg/mL)	14.24 ± 1.81	18.23 ± 3.47
<i>A. millefolium</i> (100 µg/mL)	16.91 ± 1.63 *	38.96 ± 8.35 *

(caption on next page)

**Fig. 3.** The effect of *A. millefolium* hydroethanolic extract on cell death by apoptosis of NCI-H460 and HCT-15 cells. Cells were treated for 48 h with 75  $\mu\text{g/mL}$  or 100  $\mu\text{g/mL}$  of the hydroethanolic extract, or with the corresponding volumes of water (control 1 and control 2, respectively). Panels a. and b. correspond to representative histograms of the flow cytometry analysis of apoptotic cell death following Annexin V-FITC/PI staining in NCI-H460 and HCT-15 cells, respectively. Images are representative of three independent experiments(c) Levels of programmed cell death. Results are the mean  $\pm$  SE of three independent experiments. \* $p \leq 0.05$  treatment vs the respective control.



**Fig. 4.** Levels of p53, pro-caspase 3, total PARP and p21 in NCI-H460 cells following treatment with the *A. millefolium* extract, analyzed by Western blot. Cells were treated for 48 h with complete medium (Blank), Controls (vehicle of the extract,  $\text{H}_2\text{O}$ ), or with 75  $\mu\text{g/mL}$  or 100  $\mu\text{g/mL}$  of the extract. Tubulin or actin was used as loading control. Left panel shows a representative image of the Western blots. Right panel shows the densitometry analysis expressed after normalization of the values obtained for each protein with the values obtained for actin or tubulin and representing the mean  $\pm$  SE from at least 3 independent experiments. \* $p \leq 0.05$  between treatment and Control 2.



**Fig. 5.** Levels of XIAP, pro-caspase 3 and total PARP in HCT-15 cells following treatment with *A. millefolium* extract, analyzed by Western Blot. Cells were treated for 48 h with complete medium (Blank), Controls (vehicle of the extract,  $\text{H}_2\text{O}$ ), or with 75  $\mu\text{g/mL}$  or 100  $\mu\text{g/mL}$  of the extract. Tubulin was used as loading control. A) Image representative of three independent experiments. B) Densitometry analysis expressed after normalization of the values obtained for each protein with the values obtained for tubulin and representing the mean  $\pm$  SEM from 3 independent experiments. \* $p \leq 0.05$  treatment vs Control 2.

hydroxycinnamic acid, being chlorogenic acid derivatives the major compounds present. Regarding flavonoids, 11 were identified as flavones, mainly apigenin and luteolin glycoside derivatives, and 9 as flavonols being quercetin, kaempferol and isorhamnetin glycoside derivatives the main aglycones present. These compounds were previously identified and described by Dias et al. for aqueous and methanolic extracts of the same plant species (Dias et al., 2013).

*cis* and *trans* 3,5-O-Dicaffeoylquinic acid, followed by 5-O-caffeoylquinic acid, were the main phenolic acids, accounting to 66% of the total phenolic compounds. While, luteolin-O-acetylhexoside and apigenin-O-acetylhexoside were the main flavonoids, consisting in 9% of the polyphenols content. These compounds have been previously described to have cytotoxic activities (Costa et al., 2014; In et al., 2016; Lin et al., 2015; Murad et al., 2015).

### 3.2. The *A. millefolium* hydroethanolic extract inhibited the growth of human tumor cell lines

A screening of the *A. millefolium* hydroethanolic extract was performed with the sulforhodamine B assay in two human tumor cell lines:

NCI-H460 and HCT-15. Results (Table 2) showed an inhibitory effect of this plant extract on the growth of all the tested cell lines. The  $\text{GI}_{50}$  (concentration which inhibits the cell growth of 50% of cells) obtained was 187.3 and 70.8 for NCI-H460 and HCT-15 cells, respectively. This effect was more potent in the colon cancer cell line, and secondly in the lung cancer cell line. This bioactivity could be related to the high percentage of phenolic acids, mainly chlorogenic acid derivatives, accounting to 80% of the total phenolic composition present in the extracts. Chlorogenic acids have been previously described to have antiproliferative activity against several cell lines (Costa et al., 2014; In et al., 2016; Lin et al., 2015; Murad et al., 2015). Based on this, the NCI-H460 and the HCT-15 cell lines were further studied in order to unravel the mechanism of action of this plant extract.

### 3.3. The *A. millefolium* hydroethanolic extract reduced the number of viable cells

In order to further confirm that this extract had a cell growth inhibition effect, the viable cell number was analyzed with the Trypan Blue exclusion assay in the colon and lung cancer cell lines. The number

of viable cells was reduced when cells were treated for 48 h with *A. millefolium* (75 µg/mL and 100 µg/mL). As previously observed with the SRB assay, the extract was more potent in the HCT-15 cell line. In the NCI-H460 cells, when compared with the blank cells the extract caused a reduction of 28% and 55% in the viable cell number when tested at concentrations of 75 µg/mL and 100 µg/mL, respectively (Fig. 1a). In the HCT-15 cells, the same concentrations caused a more pronounced reduction in the number of viable cells, of 75% and 82% respectively (Fig. 1b).

### 3.4. The *A. millefolium* hydroethanolic extract interfered with the cell cycle progression

In order to unravel the molecular pathways involved in the growth inhibitory effect of the *A. millefolium* extract in both the colon and lung cancer cell lines, the cell cycle profile was analyzed by flow cytometry following incubation with propidium iodide. In NCI-H460 cells, the extract (100 µg/mL) caused a statistically significant decrease of 6.0% in the G2/M phase of the cell cycle when compared to Control 2 cells (Fig. 2a). In addition, this concentration caused a slight increase (non-statistically significant) in the % of cells in the S phase of the cell cycle, when compared to control treatments.

On the other hand, treatment of HCT-15 cells with *A. millefolium* (100 µg/mL) caused a statistically significant decrease (of 11.4%) on the number of cells in the G1 phase of the cell cycle, when compared to Control 2 and a slight increase (non-statistically significant, of 4%) in the number of cells in G2/M (Fig. 2b).

### 3.5. The *A. millefolium* hydroethanolic extract induced cell death in the NCI-H460 and HCT-15 cells

The effect of this plant extract in programmed cell death by apoptosis was also analyzed in both the lung and colon cell lines, with the Annexin-V and PI protocol. The obtained results (Fig. 3) showed that this extract when used at a concentration of 100 µg/mL caused an increase in the % of cells undergoing apoptosis in both cell lines, when compared to the respective control treatments. Moreover, the effect was more pronounced in the HCT-15 cell line, which is in agreement with the cell growth and viability results previously obtained. Indeed, in the HCT-15 cell line, a lower concentration of extract (75 µg/mL) also induced apoptosis, even though this increase was not considered statistically significant.

### 3.6. The *A. millefolium* hydroethanolic extract alters the expression levels of proteins involved in cell cycle and apoptosis in the NCI-H460 and HCT-15 cells

Considering the here obtained results, showing effect on cell cycle profile and apoptosis, the expression of some proteins related to these pathways was studied in both cell lines, following treatment with 75 µg/mL and 100 µg/mL of the extract. Interestingly, the obtained results were different between the two cell lines.

In the NCI-H460 cells, results showed that treatment with the *A. millefolium* extract caused an increase in the levels of p53 and p21, and a slight decrease in the levels of pro-caspase 3 and total PARP (Fig. 4). However, this effect was only verified with the highest concentration tested (100 µg/mL) and was considered statistically significant only for p21. In addition, no statistically significant effect was seen in the expression of XIAP (even though a slight increase was detected, data not shown).

In the HCT-15 cells, results revealed that the extract caused a reduction in the expression levels of XIAP, pro-caspase 3 and total PARP (Fig. 5). These results were observed at both concentrations tested (75 µg/mL and 100 µg/mL) and were considered statistically significant for all the proteins analyzed: XIAP, pro-caspase 3 and total PARP. In addition, no effect was seen on p53 and p21 (data not shown) (see

Fig. 5).

## 4. Discussion

This work demonstrates for the first time that a hydroethanolic extract of *A. millefolium* has antitumor potential by interfering with the progression of the cell cycle and causing apoptosis, this could be due to the fact that this extract has a high concentration in chlorogenic acid derivatives (mainly 3,5-*O*-dicaffeoylquinic acid and 5-*O*-caffeoylquinic), known to have a high bioactive capacity, especially in the inhibition of cell growth (Anantharaju et al., 2016; Gouthamchandra et al., 2017). This extract, rich in chlorogenic acid derivatives, was more potent in the HCT15 cell line than in the NCI-H460 cell line, suggesting a different mechanism of action in the two cell lines, which could be an effect of polyphenol composition present in the extract. There are few studies that have also reported the therapeutic efficacy of chlorogenic acids derivatives for treating cancers, revealing that these hydroxycinnamic acid derivatives down modulate MAPK and AKT signaling pathways to inactivate the NF-κβ, AP-1 and STAT3 in lung adenocarcinomas (Tsai et al., 2013), thus phenolic acids, especially hydroxycinnamic acid derivatives are known to play a crucial role in blocking oncogenic pathways to inhibit cancer progression (Anantharaju et al., 2016). Moreover, a number of pre-clinical and phase I clinical studies have shown that treatment with chlorogenic acid derivatives have shown beneficial effects in colon, lung and brain cancer, breast tumours, and chronic myelogenous leukemia (Bandyopadhyay et al., 2004; Belkaid et al., 2006; Xu et al., 2013).

Indeed, in the two cell lines that were used (the NCI-H460 and the HCT-15), the extract interfered with distinct phases of the cell cycle progression. In addition, the extract caused a stronger effect on apoptosis in the HCT-15 cells. This could be related to the fact that the HCT-15 cell line has an heterozygous mutation in the p53 gene (Santos et al., 2013), whereas the NCI-H460 cell line has wild-type (wt) p53. The wt p53 is a well-known tumor suppressor gene that regulates several processes in cells, such as cell cycle arrest, DNA repair and apoptosis. Nevertheless, when p53 is mutated, it does not carry this role as tumor-suppressor gene (Merkel et al., 2017; Schmidt et al., 2017).

In the NCI-H460 cells in which the p53 gene is wild-type, treatment with the extract caused an increase in the levels of p53 and also of p21, a protein described as a cell cycle inhibitor (El-Deiry, 2016) and regulated directly by p53 (Vogelstein et al., 2000). In addition, in this cell line the extract caused an increase in apoptosis levels, as verified by the Annexin V/PI results and confirmed by the verified decrease in the cellular levels of pro-caspase 3 (suggesting that caspase 3 was activated (Ashkenazi and Salvesen, 2014)) and of total PARP (known to be correlated with the late stages of the apoptotic process (Duriez and Shah, 1997)). Therefore, these results suggest that the effect of the extract in this cell line is mediated via p53. This activity could be attributed to the high concentration in chlorogenic acid derivative, which may cause the increase of apoptosis levels.

In the HCT-15 cells which have mutant p53, no effect was verified in the levels of p53 or p21 upon treatment with the extract (results not shown). In addition, treatment with this extract caused a more potent effect in cell cycle and apoptosis. Nonetheless, the effect on the progression of the cell cycle was not mediated in the same way. The extract caused a decrease in the G0/G1 and a slight increase in the G2/M phases of the cell cycle, which are not compatible with a p53-mediated effect. Therefore, as expected, the effect in this cell line does not seem to involve p53. The mechanism of action of this extract in this cell line involved a very strong increase in apoptosis, as verified by the Annexin V/PI results and from the decrease in the expression levels of pro-caspase 3 and total PARP. In addition, in this cell line the extract caused a pronounced decrease in the cellular levels of XIAP. The XIAP belongs to a family of proteins named inhibitor of apoptosis proteins (IAP), which are capable of regulating several cell signaling pathways involved in numerous processes, such as cell proliferation, apoptosis,



differentiation, migration and pro-inflammatory and immune responses. Since XIAP is involved in such cellular processes, it has been largely studied as a target for new therapies, particularly in situations of resistance to apoptosis and resistance to anticancer therapy (Budhidarmo and Day, 2015; Philchenkov and Miura, 2016; Pluta et al., 2015). Indeed, various inhibitors of XIAP (either antisense oligonucleotides or small molecules) have been studied in pre-clinical and clinical trials (Flygare et al., 2012; Lee et al., 2016; Mahadevan et al., 2013). The results here presented show that this extract reduces XIAP protein levels in the HCT-15 cells, which is possibly the main mechanism of action of this extract in this cell line. These findings could be related to the main metabolites present in the extract, which are mainly constituted with chlorogenic acid derivatives. Indeed, another inhibitor of XIAP was previously documented to induce apoptosis and to arrest cell cycle in G2/M (Langemann et al., 2017).

Other studies had shown the antitumor potential of other extracts of *A. millefolium* in various human tumor cell lines. Indeed, a chloroform-soluble extract of this plant exerted tumor cell proliferation inhibitory activities on HeLa and MCF-7 cells, and also a moderate effect on A431 cells (Csupor-Löffler et al., 2009). Moreover, previous studies have shown that methanolic, infusion and decoction extracts of *A. millefolium* reduced the growth of five different cancer cell lines. In that study, the infusion extract was the most potent one, in a breast cancer cell line (MCF-7), therefore presenting the lowest GI<sub>50</sub> (Dias et al., 2013).

The chemical composition of several extracts (methanolic, infusion and decoction) from *A. millefolium* was studied and revealed that these are composed by phenolic acid derivatives, mainly caffeoylquinic acid derivatives and flavones (Dias et al., 2013). Both caffeoylquinic acid derivatives and flavones are recognized for their antioxidant, antitumor and anti-inflammatory properties (Costa et al., 2014; In et al., 2016; Lin et al., 2015; Murad et al., 2015). Therefore, the composition of different extracts (hydroalcoholic, methanolic, aqueous, infusion and decoction) suggests an antioxidant potential and this activity was demonstrated by several authors, using different assays (Candan et al., 2003; Dias et al., 2013; Kintzios et al., 2010; Vitalini et al., 2011). In addition, some sesquiterpenoids were isolated as methyl esters from *Achillea millefolium* and were active against mouse P-388 leukemia cells *in vivo* (Tozyo et al., 1994).

## 5. Conclusion

The presented results proved that the *A. millefolium* hydroethanolic extract caused reduction of cell growth in the cell lines tested. This effect was more potent in the colorectal cancer cell line (HCT-15, with mutant p53) but also had a strong effect in the lung cancer cell line (NCI-H460, with wild-type p53). This could be due to the high concentration in chlorogenic acid derivatives, which have shown that they can induce cytotoxic effects in many human cancer cell lines.

This extract, rich in chlorogenic acid derivatives, interfered with the cell cycle progression and induced apoptosis in those two cell lines. However, the mechanism of action was different in the two cell lines, possibly due to the fact that one had wild-type p53 and the other one had mutant p53. The effect in the NCI-H460 cell line was possibly mediated via p53, whereas the effect in the HCT-15 cell line was, at least in part, due to a downregulation of XIAP levels. It would be interesting to further analyze the biochemical composition of extract and to isolate the different compounds present in this extract in order to identify potent inhibitors of XIAP. Therefore, this work enhances the interest of this plant extract as a possible source of new natural bioactive compounds, particularly of XIAP inhibitors.

## Conflicts of interest

The authors declare no conflict of interest.

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