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### Defective apoptosis of U937 cells induced by benzyl isothiocyanate (BITC)

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INTRODUCTION

Isothiocyanates' precursors (ITCs), including benzyl isothiocyanate (BITC), are considered as cancer chemopreventive agents. ITC derivatives were tested in clinical trials (NCT00005883, NCT01265953, NCT01790204) and preclinical studies aimed to inhibit tumor growth and modulation of their microenvironment. Although efficacy of ITCs was demonstrated with several leukemic cell lines, the final steps of BITC-induced apoptosis were not completely elucidated in the literature. Therefore, we focused on morphological and biochemical events occurring upon treatment of U937 leukemia cells with BITC. Micromolar concentrations of BITC induced cytotoxicity in U937 cells, with major features resembling the hallmarks of apoptosis: phosphatidylserine exposure, low mitochondrial membrane potential, and presence of PARP cleavage by caspases. Disassembly to apoptotic bodies, a final step of classic apoptosis, was not observed. While tracing the signalling pathways, our results showed increased levels of BAG-1 and PUMA proteins, but in contrast to other models of ITCs-induced apoptosis, downregulation of Mcl-1 protein was not noticed. Additionally, BITC-induced dying U937 cells released lower levels of chemoattractants, such as IL-8 and MCP-1, when compared to cells undergoing classical apoptosis. This may disrupt clearance of cell debris by macrophages in vivo (efferocytosis), and in turn affect the inflammatory response. In summary, BITC inhibits tumor growth which makes it a good candidate for supporting cancer treatment. However, atypical apoptosis of leukemic U937 cells induced with BITC may affect the ability of phagocytes to effectively scavenge cellular debris, which poses a question of BITC effectiveness as a chemopreventive agent for leukemias.

Key words: benzyl isothiocyanate, apoptosis, U937, blebbing

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Chemoprevention is a strategy to inhibit, delay or reverse carcinogenesis using naturally occurring or synthetic chemical agents. Chemopreventive properties are especially attributed to isothiocyanates (ITCs) (Zhou et al., 2013), the organosulfur compounds abundantly present mainly in the cruciferous vegetables (e.g., watercress, brussels sprouts, broccoli, cabbage). ITCs are the product of enzymatic activity of a plant-specific myrosinase or glucosinolates of intestinal flora (Miyoshi et al., 2012).

Over the past three decades, research has confirmed the efficacy of various ITCs against cancer in preclinical models (Rao, 2013). One of such dietary chemopreventive agents is the benzyl isothiocyanate (BITC), which exerts anti-inflammatory properties and is characterized by the ability to inhibit growth of various chemically induced tumours (Lee et al., 2009; Rao 2013) or cancer stem cells CSC (Kim & Singh, 2015), by inducing apoptosis. BITC mediates inhiet al. bition of clonogenicity and soft-agar colony formation by breast cancer cells (Xie et al., 2017). Exposure of different cell lines to BITC promotes apoptosis with majority of its characteristic biochemical changes, e.g. caspase activation, cytochrome c release, nuclear apoptosis-inducing factor (AIF) accumulation, and Bcl2-associated X protein (Bax) translocation (Liu et al., 2013; Zhou et al., 2013). BITC is able to cause cell cycle arrest in the G2/M phase, reactive oxygen species generation and glutathione depletion (Liu et al., 2013). Molecular understanding of BITC-mediated signalling networks is still not fully elucidated (Xie et al., 2017).

Apoptosis was considered as a immunologically silent process (Negroni et al., 2015), although literature data suggests that it may promote inflammatory events which potentially recruit and activate immune cells. Aberrations of apoptosis inevitably involve adverse effects. The effectiveness of disassembly of a dying cell to apoptotic bodies should be taken under consideration in terms of cancer therapy. Such disassembly is the most critical step in apoptosis-induced immune response. Inefficient clearance of dying cells causes persistent inflammation (Wickman et al., 2013), a critical component of cancer progression. Data suggest that inflammatory microenvironment may be an indispensable participant in the neoplastic process (Bakker et al., 2016).

The study presented here was designed to determine the mechanism of BITC-induced apoptosis of leukemia cells, with special emphasis on cell disassembly. Leukemia is a white blood cells cancer, with over 54000 new cases diagnosed worldwide per year, and a 5-year survival rate below 60% (Bakker et al., 2016). We used U937 cells as a commonly utilized research model of mono-

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Abbreviations: BAG-1, BCL2-associated athanogene; PUMA, p53 Abbreviations: BAG-1, BCL2-associated atnanogene; PUIVIA, p53 upregulated modulator of apoptosis; Mcl-1, myeloid cell leu-kemia-1; CSC, cancer stem cell; PS, phosphatidylserine; VP-16, Vepesid<sup>®</sup>/etoposide; DAPI, 4',6-Diamidine-2'-phenylindole dihy-drochloride; MMP, Mitochondrial membrane potential; TMRM, te-tramethyl rhodamine methyl ester; LDH, lactate dehydrogenase; DOCK Bho associated protein kinase; DAMB, damageassociated ROCK, Rho-associated protein kinase; DAMPs, damage-associated molecular patterns

cytes. They were used in studies on cell death (Stasilojc *et al.*, 2013), novel anticancer drugs' evaluation, including ITCs action (Zhou *et al.*, 2013), as well as an *in vitro* assay to identify contact sensitizers (Python *et al.*, 2007).

Apoptosis is characterized by contraction and shrinkage of cytoplasmic organelles, as well as nuclear chromatin condensation (Wickman et al., 2013; Zhou et al., 2013). Formation of membrane blebs precedes disassembly to apoptotic bodies (Rao, 2013; Zhou et al., 2013). Exposure of phosphatidylserine (PS) on the outer leaflet of cell membrane, decrease in mitochondrial membrane potential, release of cytochrome c from mitochondria and activation of caspases (Fadok et al., 1992; Keum et al., 2004) are also the hallmarks of apoptosis. The most important final step of this process is efferocytosis followed by a "silent" immune response. Lack of pro-inflammatory activity towards cell debris after apoptosis is correlated with anti-inflammatory reaction of the neighbouring tumour and immune cells (Grabiec & Hussell, 2016). Inadequate engulfment of apoptotic bodies is followed by plasma membrane rupture and release of immunogenic intracellular materials (Grabiec & Hussell, 2016; Toda et al., 2015; Wickman et al., 2013; Zhang et al., 2018). Dysfunction of disassembly may affect the scavenging of cell remains and, indirectly, effectiveness of leukemia therapy with BITC.

Here, we report atypical morphology of BITC-treated U937 cells with significantly lower level of chemoattractants released and a decreased degree of cell disassembly to apoptotic bodies when compared to cells exposed to etoposide (VP-16), a popular anticancer chemotherapeutic drug.

#### MATERIALS AND METHODS

**Cells and Media**. U937 (cat. no. CRL-1593.2<sup>TM</sup>), Raji (cat. no. CCL-86<sup>TM</sup>) and Namalwa (cat. no. CRL-1432<sup>TM</sup>) cell lines which were derived from histiocytic lymphoma or Burkitt's lymphoma cells were obtained from ATCC<sup>®</sup>, USA. Cells were cultured at 37°C in a humidified atmosphere (5% CO<sub>2</sub>) in the RPMI 1640 medium with 2 mM L-glutamine (Cytogen, Germany), supplemented with 10% fetal bovine serum (Cytogen, Germany), 1 mM sodium pyruvate, 100 U/ml penicillin, 100  $\mu$ g/ml streptomycin. The cell lines were mycoplasma-free as tested by DAPI staining and fluorescence microscopy analysis.

Determination of BITC cytotoxicity towards U937 cells.  $1 \times 10^5$  cells were seeded in the wells of a 96-well plate, in 100 µl of the medium. The cells were treated with different concentrations of BITC (0–62.5 µM) for 24 h. 2 h before the end of the treatment, 12 µl of MTT (Sigma Aldrich, 5 mg/ml) were added to all wells. After incubation, the formazan crystals formed by viable cells were dissolved in 110 µl of acidic (40 mM HCl) isopropanol and analysed in a microplate reader at  $\lambda$ =540 nm. Absorbance values of the wells containing treated cells were compared with those of wells with non-treated cells.

Induction of cell death.  $1 \times 10^5$  of U937 cells per well of a 96-well plate or  $3 \times 10^6$  of U937, Raji or Namalwa cells per well of a 6-well plate were exposed to 25  $\mu$ M BITC (Sigma-Aldrich, 98% purity) or 50  $\mu$ g/ml VP-16 (Vepesid<sup>®</sup>, Bristol-Myers Squibb S.p.A., Italy) for up to 24h, with time range dependent on the type of analysis. After treatment, cells seeded in 6-well plates were used for morphological analysis under Axiovert 200 microscope, with 40× long distance objective. Western blot analysis. PARP cleavage, anti- and proapoptotic proteins' expression. After treatment, the cells were harvested and centrifuged (5 min,  $400 \times g$ , 4°C). Afterwards, the cells were washed twice with PBS×1 and suspended in 300 µl of Laemmli buffer (SDS 10%; Tris 0.5 M pH 6.8; glycerol 20%). Cell lysates were boiled for 5 min and cooled on ice for 3 min, with boiling and cooling processes repeated three times. Protein concentration was analysed with the NanoDrop spectrophotometer and standardised in all samples to 1 mg/ml. Samples were mixed with the modified Laemmli buffer (with addition of  $\beta$ -mercaptoethanol and bromophenol blue) and boiled for 5 min.

Cell lysate was separated on a 4% (stacking gel) and 12% (resolving gel) SDS-PAGE under reducing conditions. Resolved proteins were transferred onto a PVDF membrane which was then probed with primary antibodies: monoclonal, rabbit anti-human PARP (Cell Signalling, USA, cat no. 9532, 1:1000 dilution); monoclonal, mouse anti-human Mcl-1 (Santa Cruz Biotechnology, USA, cat. no. sc-12756, 1:200 dilution); monoclonal, mouse anti-human Bcl-2 (BD Transduction Laboratories, USA, cat no. 610538, 1:1000 dilution); monoclonal, mouse anti-human BAG-1 (BD Transduction Laboratories, USA, cat no. 611868, 1:30000 dilution); monoclonal, mouse anti-human PUMA (Cell Signalling, USA, cat no. 4976, 1:1000 dilution); monoclonal, mouse antihuman Bid (BD Transduction Laboratories, USA, cat no. 611528, 1:500 dilution); monoclonal, mouse anti-human Bad (BD Transduction Laboratories, USA, cat no. 610391, 1:1000 dilution); monoclonal, mouse anti-human β-tubulin (Santa Cruz Biotechnology, USA, cat. no. sc-55529, 1:200 dilution).

Affinity purified monoclonal goat anti-mouse IgG or goat anti-rabbit IgG (Santa Cruz Biotechnology, USA, cat no. sc-2004 and cat no. sc-2005 respectively) conjugated to HRP (horseradish peroxidase 1:2000 dilution) were used as secondary antibodies. BM Chemiluminescence Blotting Substrate (Roche, Switzerland) was used to visualize the protein bands.

Densitometric and statistical analyses for the obtained blots were performed for at least three independent repeats, with the use of Image J v 1.48 (NIH), and Statistica 12.

Flow cytometry analysis. Early apoptosis was evaluated by measuring the exposure of phosphatidylserine on the outer leaflet of plasma membrane. Analysis was performed using FITC-conjugated Annexin-V (FITC Annexin V Apoptosis Detection Kit, BD Pharmingen<sup>TM</sup>).  $1 \times 10^5$  BITC-treated cells were washed with 200 µl PBS, centrifuged and then suspended in 100 µl binding buffer. Afterwards, the cells were incubated for 5 min at 37°C with 5 µl of FITC-conjugated Annexin-V and 5 µl of propidium iodide. Four subpopulations were identified according to their fluorescence: PI-low/FITC-low (live cells), PI-high/FITC-low (necrotic cells), PI-low/FITChigh (early apoptotic cells), PI-high/FITC-high (late apoptotic cells – confirmed by kinetics of the process).

The mitochondrial membrane potential (MMP) was evaluated using tetramethyl rhodamine methyl ester (TMRM, Fluka). U937 cells were stained with 0.1  $\mu$ M TMRM for 30 min at 37°C before the end of treatment with apoptotic inducers. Afterwards, the cells were centrifuged and suspended in 250  $\mu$ l of fresh RPMI 1640 medium for analysis. Cells with lower mitochondrial potential showed a shift in fluorescence intensity from the control region towards the lower values. All of the measurements were performed for 10 000 cells that exhibited a typical forward and side scatter features of non- disassembled cells. Data were analysed off-line using the BD FACSDiva software (BD Biosciences, USA).

LDH (lactate dehydrogenase) level determination.  $1 \times 10^5$  cells were treated with 50 µg/ml VP-16 or 25 µM BITC for 5 h and supernatants were collected by centrifugation ( $400 \times g$ , 5 min). Supernatant (100 µl) was placed in a 96-well plate and incubated with 100 µl of mixed catalyst and Dye Solution (LDH Cytotoxicity Detection Kit, Clontech, USA). Additional controls were performed as follows: medium without cells (absorbance value was subtracted from all controls and samples), non-treated cells' supernatant (low control), and triton X-100 0.1% treated cells' supernatant (high control). The absorbance was analysed with a microplate reader (450 nm) after 15 min of incubation.

**Determination of cytokine concentration**.  $1 \times 10^6$  of U937 cells were treated with BITC (25  $\mu$ M) or VP-16 (50  $\mu$ g/ml) for 1.5 h, and then the cells were centrifuged (5 min, 400×g RT). Pelleted cells were resuspended in the medium devoid of cell death inducers. After 22.5 h incubation at 37°C with 5% of CO<sub>2</sub>, the cells were centrifuged. Supernatants were collected and stored at  $-70^{\circ}$ C. IL-8 and MCP-1 levels were assessed in supernatants by enzyme-linked immunosorbent assay using BD OptEIA<sup>TM</sup> Human ELISA Kits II (BD Biosciences, USA).

**Statistical analysis.** Statistical differences between diversely treated U937 cell cultures were determined using the *t*-test and U Mann-Whitney test after confirming the normal distribution of data. The analyses were performed using 3 to 4 replicate runs in at least three independent experiments. *p*-values <0.05 were considered to indicate statistically significant differences.

### RESULTS

# Atypical morphology of U937 cells treated with BITC as a cell death inducer

We analysed morphological changes of U937 cells in response to different concentrations of BITC. According to the literature, an effective concentration of BITC ranges from 0.89  $\mu$ M to 100  $\mu$ M (Lee *et al.*, 2012). Interestingly, for all tested concentrations of BITC, similar morphological changes and a negligible presence of apoptotic bodies were observed (Fig. 1a). Under our experimental conditions an effective BITC concentration which induced a decrease in cell viability after 24 h treatment was 12  $\mu$ M (Fig. 1b). However, higher concentration of BITC (25  $\mu$ M) was chosen for further investigation in order to assess the final steps of cell death in the majority of cells.

When well-known inducer of apoptosis VP-16 (etoposide) (Jafarlou *et al.*, 2016) was applied at a concentration of 50  $\mu$ g/ml, apoptotic morphological markers and apoptotic bodies were already observed after 3 h of treatment.

Application of BITC ( $25 \mu$ M) for the first 5 h of treatment did not exert a significant morphological change, in contrast to 3 h VP-16 treatment (Fig. 1c).

These results indicate that even high concentration of BITC (62.5  $\mu$ M) is not able to cause cell disassembly to apoptotic bodies, although lower BITC concentration, i.e. 12  $\mu$ M, causes cell death. Unlike treatment with VP-16, BITC did not cause *blebbing* and disassembly of U937 cells.



Figure 1. Cytotoxicity of BITC and morphology of the dying U937 cells.

(a) Light microscopy photographs of cells treated with BITC (0–62.5  $\mu$ M) for 24 h (40× objective). (b) Viability of U937 cells (MTT assay) after 24 h treatment with different concentrations of BITC (0–30  $\mu$ M). Results show changes in metabolic activity of cells (in percentage)  $\pm$  S.D. (n=4). (c) Light microscopy photographs of cells treated with 25  $\mu$ M of BITC or 50  $\mu$ g/ml VP-16 for 24 h (40× objective).

## Identification of the type of cell death of BITC-treated U937 cells

BITC was unable to induce morphological markers of apoptosis in dying U937 cells, which remains in discordance with previous reports that suggests that BITC induces this type of cell death (Stasilojc *et al.*, 2013; Wu *et al.*, 2011). Therefore, we decided to analyze other markers of apoptosis as well.

One of the most characteristic markers of the ongoing apoptosis is translocation of phosphatidylserine to the outer leaflet of cell membrane. Presence of phosphatidylserine on BITC-treated U937 cells was analysed with flow cytometry. Simultaneous staining of cells with propidium iodide was used to distinguish necrotic cells. Our results confirmed translocation of phosphatidylserine in BITC-treated cells, however, this process was delayed in comparison to the VP-16-treated cells (Fig. 2a).

Dying U937 cells were characterized by the decreased mitochondrial membrane potential (MMP). The obtained results were similar for both, the BITC- and VP-16-treated U937 cells (Fig. 2b).

To define if the observed cell death mechanism is caspase-dependent, presence of cleaved PARP in cell lysates was evaluated. Cleaved form of PARP was pre-



#### Figure 2. Apoptotic features of U937 cell death.

(a) Phosphatidylserine presence on the outer leaflet of cell membrane of U937 cells treated with 25  $\mu$ M BITC or 50  $\mu$ g/ml VP-16 for 0–4 h. Cells analysed by flow cytometry after annexin-V-FITC and propidium iodide staining. Results show the mean number of cells with positive FITC signal (in percentage of all analysed cells)  $\pm$ S.D. (*n*=4). Statistically significant differences between cells treated with BITC and VP-16 are indicated (\**p*<0.05). (b) Mitochondrial membrane potential (MMP). Cytometric analysis of MMP with TMRM in cells treated with 25  $\mu$ M BITC or 50  $\mu$ g/ml VP-16 for 4 h. Results represent the mean number of cells with high MMP (high TMRM – considered as the level of fluorescence for control cells)  $\pm$ S.D. (*n*=4). Statistically significant differences between U937 cells treated with BITC or VP-16 and control are indicated (\**p*<0.05, \*\**p*<0.005). (c) PARP cleavage. U937 cells treated with 25  $\mu$ M BITC for 0–5 h. Cytosolic fractions of cell lysates were prepared and subjected to western blot analysis using antibodies against PARP. Presence of 116 kDa PARP protein and one of its cleaved fragments, 85 kDa CF (catalytic fragment) were visualised. β-tubulin was used as a protein quantity marker. (d) LDH release. Cells were treated with 25  $\mu$ M BITC or 50  $\mu$ g/ml VP-16 for 0–5 h. The cell supernatants were obtained and analysed in colorimetric test for the presence of cytoplasmic LDH (lactate dehydrogenase). Results show the mean value of LDH activity as percentage ± S.D. (*n*=5) of maximal absorbance control (high control). High absorbance indicates a presence of cells with discontinuous cell membrane). High control – U937 cells with discontinuous cell membrane).

sent in cell lysates after 2 h of treatment and its amount increased in a time dependent manner (Fig. 2c).

Additionally, we looked for a possible necrotic component of the observed cell death (Miyoshi *et al.*, 2008). The level of lactate dehydrogenase (LDH) released from cytosol was assessed in supernatants of cells treated with BITC or VP-16 for 5 h. Activities of LDH in the supernatants of treated and untreated cells were compared. The results showed a similar, low level of LDH release for both tested cell death inducers, which indicated a lack of rapid necrosis (Fig. 2d). In summary, U937 cells treated with BITC exhibit

In summary, U937 cells treated with BITC exhibit most of the characteristics of caspase-dependent apoptotic cell death, similar to those observed for the U937 cells treated with VP-16, except for disassembly to apoptotic bodies which is a morphological marker of late apoptosis.

#### BITC-treatment does not affect Bcl-2, Mcl-1, Bid and Bad protein levels, but affects PUMA and BAG-1 in the U937 cells

The main regulators of apoptosis are the Bcl-2 family of pro- and anti-apoptotic proteins. A dependence between the level of myeloid cell leukemia-1 (Mcl-1 – BCL2 Family Apoptosis Regulator) and BITC-induced apoptosis was suggested (Jafarlou *et al.*, 2016; Zhou *et al.*, 2013). During the first 12 h of BITC treatment, no statistically relevant changes in the Mcl-1 level were found in the U937 cells. Similar result was obtained for major anti-apoptotic protein Bcl-2. However, we observed a statistically significant increase in the levels of large and medium isoforms of BAG-1 (BCL2-associated athanogene) during the first 6 hours of treatment. BAG-1 small isoform showed a distinct trend towards significance (p<0.07) after 12 h of BITC-treatment (Fig. 3). Densitometric analysis is shown in the supplementary data.



Figure 3. Level of pro- and anti-apoptotic proteins in cell lysates.

Lysates were obtained from U937 cells treated with 25  $\mu$ M BITC for 0–6 and 12 h. The expression of McI-1, BcI-2, BAG-1, PUMA, Bid, Bad was analysed with Western blot.  $\beta$ -tubulin was used as a protein quantity marker.



Figure 4. Cytokine release by dying U937 cells.

U937 cells were induced to die with 25  $\mu$ M of BITC or VP-16 (50  $\mu$ g/ml) for 1.5 h followed by a change of medium. The cell supernatants were obtained after 24h and analysed by ELISA test. (a) IL-8 concentration. (b) MCP-1 concentration. Results represent the concentration of cytokines (pg/ml) in the supernatants ±S.D. (n=3). Statistically significant differences between U937 cells treated with BITC or VP-16 indicated (\*\*p<0.01, \*\*\*p<0.005), between inducer-treated cells and untreated cells indicated (\*p<0.5, \*\*p<0.01, \*\*\*p<0.005)

Increased expression of the PUMA protein (p53 upregulated modulator of apoptosis) in adherent cell lines treated with BITC was indicated in the literature (Antony *et al.*, 2012) and our results remain in accordance with that report. A significant increase in the PUMA level was observed during BITC-treatment, whereas the Bid and Bad protein levels did not changed during the first 12 h of treatment (Fig. 3).

#### Release of chemotactic cytokines

IL-8 and MCP-1 are two of the most important proinflammatory chemokines (Hu *et al.*, 2014). MCP-1 recruits monocytes into the foci of active inflammation and it is associated with the development of Th2-polarized response (Deshmane *et al.*, 2009). Supernatants collected from BITC- or VP-16-treated cells (24 h) were used to determine the level of IL-8 and MCP-1.

Unlike the control and VP-16-treated cells, BITCtreated U937 cells secreted a significantly lower level of IL-8 and MCP-1 (Fig. 4a and Fig. 4b). Thus, it should be emphasized that BITC-treated U937 cells, which did not disassemble during the death process, secrete lower levels of chemotactic cytokines.

## Presence of that unusual cell death morphology in other cell lines

Raji and Namalwa cells were treated with BITC (25  $\mu$ M) or VP-16 (50  $\mu$ g/ml) for 24h. Those cells did show similar morphological changes to those of BITC-treated U937 cells (Fig. 5). However, morphological dif-



Figure 5. Morphology of dying Raji, Namalwa and Ramos cells. Light microscopy photographs of cells treated with BITC (25  $\mu$ M) or VP-16 (50  $\mu$ g/ml) for 24 h (40× objective).

ferences between cells treated with BITC and VP-16 were not as evident as for the U937 cells.

#### DISCUSSION

Few reports describe cell death of human or murine leukemia cells (Jurkat, HL-60, WEHI-3) treated with BITC (Antony et al., 2012; Coleman et al., 2001; Lee et al., 2009; Zhou et al., 2013). In the study presented here we chose U937 cells as an acknowledged model for a standard caspase-dependent apoptosis (Stasilojc et al., 2013). Our results remain in accordance with previous reports of BITC cytotoxic effects on leukemia cells (Zhang et al., 2003), but indicate that the observed morphological and biochemical markers are not typical for standard apoptotic cell death (Fadok et al., 1992). Pharmacokinetic data for BITC in the human serum is currently unavailable. However, in vivo concentrations of BITC used in animal studies are several orders of magnitude higher (e.g. 10 mg/kg equal to 50  $\mu$ M) than dietary consumption in humans (Kassie et al., 1999). In our cell model (U937) 25 µM of BITC was close to IC90, so it was possible to observe and describe changes induced by BITC in a late stage of the apoptosis process.

One of our main goals was to observe morphological effects of BITC on U937. Negligible cell disassembly, but noticeable granularity of the dying cells were observed. These morphological markers were different in comparison to leukemia cells treated with other death inducers (Stasilojc *et al.*, 2013). Similar morphological changes as BITC-treated U937 cells were observed for Raji and Namalwa cell lines. However, morphological differences of BITC- or VP-16-treated cells were not as evident as for the U937 cells.

Increased activity of caspases followed by cleavage of PARP (Fig. 2c), phosphatidylserine translocation (Fig. 2a) and decrease in mitochondrial membrane potential (MMP) (Fig. 2b) observed in BITC-treated U937 cells are consistent with literature description of apoptosis (Miyoshi et al., 2008; Wu et al., 2011; Xiao et al., 2006). We did not confirm the presence of a necrotic component of cell death during the first five hours after cell death induction (Fig. 2d), although such an observation was described in the literature (Miyoshi et al., 2008). Presence of secondary necrosis was negligible, as the average ratio of early apoptosis versus late apoptosis was 47.2 (±14.3). Therefore, cell death induced by BITC in U937 seems to demonstrate the main molecular markers of caspase-dependent apoptosis. However, the lack of apoptotic bodies formation indicates an atypical process of apoptosis.

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Figure 6. A proposed mechanism for the benzyl isothiocyanate (BITC)-induced U937 apoptosis with no blebbing and disassembly to apoptotic bodies.

Literature data describe BITC-induced molecular markers associated with cell death e.g. Mcl-1 or Bax downregulation, using different cell lines (Xiao et al., 2006; Zhou et al., 2013). However, our results with the U937 cells show differences with respect to these proteins. Moreover, there are reports presenting a critical role of pro-apoptotic PUMA protein in the BITC-dependent apoptosis (Antony et al., 2012). Our observations confirm the involvement of PUMA and also of the BAG-1 isoforms in the U937 model. BITC increases phosphorylation of extracellular signal-regulated kinase (ERK) (Xie et al., 2017). BAG-1 is up-regulated in a MAPK/ERKdependent fashion (Li et al., 2015). Overexpression of Mcl-1, an anti-apoptotic Bcl-2 family member protein, is associated with tumor progression and drug resistance in leukemia and several cancers (Wu et al., 2011). Literature data also indicate that apoptosis of BITC-treated leukemia cells may depend on translational downregulation of Mcl-1 (Zhou et al., 2013). Notably, the role of Mcl-1 in the death process of BITC-treated U937 cells was not indicated by our results. BAG-1 upregulation is considered as an important biomarker of increased cell survival capacities. It is also co-involved in delayed cell apoptosis activation by protection of Bcl2 and Mcl-1 proteins from degradation (Aveic et al., 2015). Insignificant changes in the Mcl-1 and Bcl2 levels may be the results of high BAG-1 level during BITC-induced cell death. On the other hand, they may also clarify unusual apoptosis activation and progression.

Disassembly of cells to apoptotic bodies followed by their clearance (efferocytosis) is one of the most important events for the immune system (Grabiec & Hussell, 2016). Membrane blebs function as chemoattractants, recruiting monocytes to the sites of apoptosis (Segundo *et al.*, 1999). The DFNA5 protein is considered as a central regulatory protein for disassembly and progression of apoptotic cells to secondary necrosis (Rogers *et al.*, 2017). We do not exclude participation of DFNA5 cleavage form in the analysed mechanism, however, the insignificant level of secondary necrosis in U937 cells in the first key time points of BITC induction suggests that its role in that mechanism may be negligible.

ROCK (Rho-associated protein kinase) plays a key role in actin-myosin contraction in bleb production. ROCK inhibition significantly reduces apoptotic body formation and apoptotic cell phagocytosis (Wickman *et al.*, 2013). Raf-1 kinase is able to reduce activity of ROCK (Matsubara & Bissell, 2016), but Raf-1 ability to inhibit ROCK depends on BAG-1 (Planchamp *et al.*, 2008). Our results showed increasing levels of BAG-1 isoforms. That result may provide a novel mechanism to explain the lack of BITC-treated U937 cell *blebbing* and disassembly to apoptotic bodies (Fig. 6).

Higher number of non-disassembled apoptotic cell remains could possibly influence the efficiency of the clearance process. Weakened efferocytosis may cause secondary necrosis of the cell debris (Rogers *et al.*, 2017). That process is followed by release of intracellular contents, including damage-associated molecular patterns (DAMPs), resulting in an inflammatory response of the innate immune system (Grabiec & Hussell, 2016; Rogers *et al.*, 2017; Wickman *et al.*, 2013).

An anti-inflammatory effect of BITC was shown with mouse cells and tissues (Lee *et al.*, 2009). Our study indicates low levels of chemoattractants (IL-8 and MCP-1) in cell supernatants. Therefore, inadequate chemoattraction of macrophages, neutrophils and human T lymphocytes towards dying U937 cells may take place (de Boer *et al.*, 2000).

### CONCLUSIONS

The BITC-treated cells were characterized by major morphological and biochemical markers of apoptosis. The imbalance between pro- and antiapoptotic proteins may influence disassembly into the apoptotic bodies. Relationship between proapoptotic PUMA and antiapoptotic BAG-1, influenced by other proteins such a Raf-1, could affect bleb formation and lack of disassembly to apoptotic bodies.

Low levels of chemokines may contribute to reduced chemotaxis of macrophages and other immune cells. Therefore, it may affect their ability to remove remnants of the dying BITC-treated U937 cells. The size of remnants and potentially lower number of recruited phagocytes may result in secondary necrosis of the dying cells *in vivo*. Tumor cells are able to suppress immune response *in vivo* (Munn & Bronte, 2016) and this effect may also be relevant in case of leukemias (Bakker *et al.*, 2016). Importantly, release of intracellular contents, including DAMPs, followed by an inflammatory response may disrupt immunosuppressive potential of the tumor cells.

Regulation of disassembly to apoptotic bodies of BITC-treated cells should be further analysed. It should cover caspase activity during that process, confirmation of ROCK involvement by using siRNA to knock it down at different time points, and finally verification of the clearance process using macrophages (e.g. THP-1) *in vitro*.

Analysis of BITC in clinical trials should also take under consideration the combined effect of defective apoptosis and decreased expression of pro-inflammatory cytokines in the tumour cells on its chemopreventive role *in vivo*. We also conclude that BITC inclusion in anticancer therapies may affect the proinflammatory effect for various cancer cell types that exhibit the same BITC-induced apoptosis mechanism as was shown for the U937 cells.

### **Conflict of interest**

We declare no conflict of interest.

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