New aspects of antiproliferative activity of 4-hydroxybenzyl isothiocyanate, a natural H₂S-donor

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Abstract
The effect of 4-hydroxybenzyl isothiocyanate (HBITC), a natural H₂S-donor from white mustard seeds (Sinapis alba), on the proliferation of human neuroblastoma (SH-SY5Y) and glioblastoma (U87MG) cells was studied and some aspects of the mechanism of its activity were suggested. The inhibition of both SH-SY5Y and U87MG cell proliferation was associated with an increase in the thiosulfate level, the number of cells with the inactive form of Bcl-2 protein, and with a decrease of mitochondrial membrane potential. Interestingly, HBITC results in downregulation of p53 protein and upregulation of p21 protein levels in SH-SY5Y cells. In the presence of elevated levels of H₂S and thiosulfate, the sulfhydryl groups of p53 protein as well as Bcl-2 protein could be modified via HBITC-induced S-sulfuration or by oxidative stress. It seems that the induction of p21 protein level is mediated in SH-SY5Y cells by p53-independent mechanisms. In addition, HBITC-treatment caused downregulation of the level of mitochondrial rhodanese and 3-mercaptopyruvate sulfurtransferase, and consequently increased the level of the reactive oxygen species in SH-SY5Y cells.

Keywords 4-Hydroxybenzyl isothiocyanate · Hydrogen sulfide · Thiosulfate · Apoptosis · Sulfurtransferases · Cancer cells

Abbreviations
SH-SY5Y The human neuroblastoma cell line
U87MG The human glioblastoma cell line
HBITC 4-Hydroxybenzyl isothiocyanate
TST Rhodanese
CTH Cystathionine-γ-lyase
CBS Cystathionine-β-synthase
MPST 3-Mercaptopyruvate sulfurtransferase
Bcl-2 B-cell lymphoma-2
MMP Mitochondrial membrane potential
H₂S Hydrogen sulfide
GSH Glutathione persulfide
SQR Sulfide:quinone oxidoreductase
GSH Reduced glutathione
ROS Reactive oxygen species
PPAR-γ Peroxisome proliferator-activated receptor-γ
LDH Lactate dehydrogenase
Sp1 Specificity protein 1
AP-2 Activating protein-2
E2Fs Elongation 2 factors
STATs Signal transducers and activators of transcription

Introduction
Hydrogen sulfide (H₂S) is endogenously generated in mammalian cells via enzymatic and non-enzymatic pathways. The enzymatic pathways generate H₂S from l-cysteine and L-homocysteine using cystathionine-β-synthase (CBS), cystathionine-γ-lyase (CTH), and cysteine aminotransferase with 3-mercaptopyruvate sulfurtransferase (MPST). H₂S can be also produced from d-cysteine by MPST with an earlier conversion by D-amino acid oxidase. Non-enzymatic production of H₂S can occur through glutathione-dependent conversations (Szabo and Papapetropoulos 2017; Tkacheva et al. 2017; Rose et al. 2017; Yagdi et al. 2016; Zheng et al. 2015; Zhang et al. 2017a). H₂S is oxidized by sulfide quinone oxidoreductase (SQR) into a sulfane sulfur atom, which is next transferred to sulfite forming thiosulfate. Sulfane sulfur atoms...
from thiosulfate can be transferred to reduced glutathione (GSH) by rhodanese (thiosulfate sulfurtransferase, TST) generating glutathione persulfide (GS-SH). Mitochondrial sulfur dioxygenase converts the sulfane sulfur persulfide forming sulfite. Then, sulfite is oxidized to sulfate by sulfite oxidase or is degraded by SQR to produce thiosulfate (Yagdi et al. 2016; Kabil et al. 2014; Jurkowska et al. 2014).

In the human body, natural hydrogen sulfide donors, releasing H2S non-enzymatically or enzymatically, include organic polysulfides from garlic (Allium sativum) or isothiocyanates from brassicas (Brassicaceae) (Tkacheva et al. 2017; Szabo and Papapetropoulos 2017). These natural products have shown interesting properties, such as antimicrobial, anti-thrombotic, anti-oxidant and anti-tumor (Song et al. 2014; Gupta et al. 2014a; Dufour et al. 2015). However, some of these effects of H2S, i.e. the anti-cancer potential, remain controversial and some of them depend on its concentration (Baskar and Bian 2011; Hellmich et al. 2015; Yagdi et al. 2016).

Citi et al. (2014) for the first time reported that some naturally occurring isothiocyanates (allyl isothiocyanate, 4-hydroxybenzyl isothiocyanate, benzyl isothiocyanate, and erucin) behave as a slow H2S-releasing agent. Among these compounds, 4-hydroxybenzyl isothiocyanate (HBITC), a natural product obtained from white mustard seeds (Sinapis alba), was the most effective, exhibiting significant and remarkable H2S-release, both in the absence and in the presence of l-cysteine (Citi et al. 2014).

Based on these data, we decided to choose HBITC to investigate the effects and mechanisms of its action in the regulation of human neuroblastoma (SH-SY5Y) and glioblastoma (U87MG) cell proliferation. It was previously demonstrated by Jurkowska et al. (2017) that diallyl trisulfide, a garlic-compound, could inhibit proliferation of these cells by its effect on Bcl-2 protein. The inhibition of U87MG cell proliferation was correlated with an increased level of sulfane sulfur (in these cells, the level of l-cysteine is higher than in SH-SY5Y cells) (Jurkowska et al. 2017).

In the present study, it was shown for the first time that HBITC, a natural H2S-donor, could reduce proliferation of both SH-SY5Y and U87MG cells. This effect was correlated with the increased level of thiosulfate, so it seems that thiosulfate as a product of H2S oxidation, and on the other hand a donor of sulfane sulfur, can be involved in the mechanisms that lead to inhibition of cell proliferation. The antiproliferative activity of HBITC was also associated with downregulation of mitochondrial rhodanese and MPST protein levels, a loss of mitochondrial membrane potential, inactivation of Bcl-2 protein and modulation (S-sulfuration, oxidative stress) of proteins involved in the cell cycle.

### Materials and methods

#### Sources of chemicals

Crystal violet (N-hexamethylpararosaniline), hexadecyltrimethylammonium bromide (CTAB) and albumin were obtained from Sigma-Aldrich Corp. (St. Louis, MO, USA). Trypsin, and penicillin/streptomycin were obtained from HyClone Laboratories (Utah, USA). The Cytotoxicity Detection Kit (LDH) and the Cell Proliferation ELISA, BrdU (colorimetric) Kit were obtained from Roche Applied Science (Germany). Potassium cyanide (KCN) was obtained from Merck (Darmstadt, Germany). All the other chemicals were of reagent grade and purchased from common commercial suppliers.

### Cell culture

Human SH-SY5Y and U87MG cells (ECACC, UK) were grown in Dulbecco’s Modified Eagle’s (DMEM) medium (Biosera, France) containing stable l-glutamine, 4500 mg/l glucose and sodium pyruvate, to which 10% fetal bovine serum (FBS) (Biowest, South America) and 1% penicillin/streptomycin (100 Units/ml penicillin and 100 µg/ml streptomycin) were added. The cells were maintained at 37 °C in a humidified atmosphere containing 5% CO2.

4-Hydroxybenzyl isothiocyanate (Santa Cruz Biotechnology, Texas, USA) was dissolved in dimethyl sulphoxide (DMSO; Sigma-Aldrich Corp., St. Louis, MO, USA), and then diluted with DMEM to the desired concentration prior to its use. The final concentration of DMSO in the medium was less than 0.1%.

### Determination of HBITC cytotoxicity

Cell cytotoxicity was investigated by measuring the leakage of lactate dehydrogenase (LDH) from dead or dying cells using the Cytotoxicity Detection Kit (Roche Applied Science, Germany) according to the manufacturer’s protocol as described previously (Jurkowska et al. 2011). The concentration of HBITC (20 µM, 40 µM, 60 µM, 80 µM) that yielded LDH leakage of less than 5% was used in the experiments.

### Cell proliferation

For the determination of cellular proliferation, the cells were seeded on 96-well plates at a concentration of 1.5 × 10^3 cells/well (SH-SY5Y cells) and 1.2 × 10^3 cells/well (U87MG cells) in DMEM supplemented as reported above. Following 24 h of incubation, the culture medium was replaced with 100 µl of complete medium with DMSO (as the controls) or...
100 µl of the medium containing various HBITC concentrations (20, 40, 60, 80 µM), and then the cells were cultured for 24 and 48 h. Cell proliferation was examined using the Cell Proliferation ELISA, BrdU (colorimetric) test (Roche Applied Science, Germany) according to the manufacturer’s protocol and also the modified crystal violet staining method (Gillies et al. 1986) was used. The absorbance was measured using an Epoch Microplate Spectrophotometer (BioTek Instruments Inc, VT, USA).

Detection of H$_2$S in cells

A H$_2$S fluorescent probe WSP-5 (Cayman Chemical, Michigan, USA) was used for detection of H$_2$S released from HBITC, according to the method of Peng et al. (2014). SH-SY5Y and U87MG cells were cultured with HBITC (40, 60, 80 µM) or the same volume of DMSO in the medium (as the controls). Then, the cells were incubated with a H$_2$S fluorescence probe WSP-5 (50 µM) and a surfactant hexadecyltrimethylammonium bromide (CTAB, 100 µM) in PBS at 37 °C for 20 min in the dark. After the PBS was removed, H$_2$S-derived fluorescence was observed under a fluorescent microscope (Axioplan-2; Carl Zeiss Imaging, Oberkochen, Germany).

Measurement of thiosulfate

For the measurement of thiosulfate, the samples of the culture medium were deproteinized using Amicon Ultra 0.5-mL centrifugal filters with a 3 kDa cutoff (Millipore, Merck, Poland), centrifuged at 14 000 x g for 20 min and assayed for thiosulfate by the method of Shih et al. (1979).

Western blot

The cells were plated at a density of 2.5 × 10$^4$ per well in six-well plates. On the following day, the cells were treated with various concentrations of HBITC or an equal volume of DMSO in the culture medium for 24 and 48 h. Proteins from the cells were extracted in lysis buffer containing 50 mM Tris–HCl, pH 7.5, 150 mM NaCl, 1 mM EDTA, 0.5% NP-40, and 1X Complete Protease Inhibitor Cocktail (Sigma-Aldrich Corp., St. Louis, MO, USA). Cell lysates were centrifuged at 20 000 x g for 15 min at 4 °C. Protein concentration was determined by the bicinechonic acid (BCA) method using a BCA Protein Assay Kit (Thermo Scientific/Pierce Biotechnology, Rockford, IL, USA) with bovine serum albumin (BSA) as a standard. The samples (25 µg) were subjected to SDS-PAGE (12%, w/v, polyacrylamide) and the separated proteins were transferred onto polyvinylidene difluoride (PVDF, Bio-Rad, CA, USA) membrane followed by blocking of the membrane with 5% nonfat milk. The sources and dilutions of primary antibodies were as follows: anti-TST (1:800; Proteintech Group, USA), anti-MPST (1:800; GeneTex, Inc., CA, USA), anti-CTH (1:1000; Abnova, Taiwan), anti-p53 (1:800; Upstate & Chemicon, CA, USA), anti-p21 (1:800; Upstate & Chemicon, CA, USA), anti-β-actin (1:1000; Sigma-Aldrich Corp., St. Louis, MO, USA). The membranes were incubated with anti-mouse or anti-rabbit goat antibodies conjugated with alkaline phosphatase (1:2000, Proteintech Group, USA) and the immune complexes were visualized using the NBT/BCIP Stock Solution (Roche Applied Science, Germany). The densities of the bands were quantified using the ChemiDocTM MP Imaging System (Bio-Rad, USA). β-actin was used as the internal control.

Bcl-2 Expression assay

Bcl-2 expression was analyzed using the Muse™ Bcl-2 Activation Dual Detection Kit (Millipore, Billerica, MA, USA). The assay utilizes two directly conjugated antibodies, a phospho-specific anti-phospho-Bcl-2 (Ser70)-Alexa Fluor 555 and an anti-Bcl-2-PECy5 conjugated antibody to measure total levels of Bcl-2 expression. The cells were prepared for this analysis as described previously (Jurkowska et al. 2017) and were analyzed by a Muse™ Cell Analyzer and a Muse™ analysis software (Merck Millipore, USA).

Oxidative stress assay

The quantitative measurement of cellular populations undergoing oxidative stress was performed using the Muse Oxidative Stress Kit (Merck Millipore, Billerica, MA, USA) according to the manufacturer’s instructions. This assay utilizes dihydroethidium (DHE), which is cell membrane-permeable and upon reaction with superoxide anions undergoes oxidation to form DNA-binding fluorophore. The kit determines the percentage of cells that are negative [ROS(−)] and positive [ROS(+)] for reactive oxygen species. Briefly, 1 × 10$^6$ cells/ml were harvested, washed with PBS, and then incubated in the dark at 37 °C for 30 min with the Muse Oxidative Stress Reagent working solution, which contained DHE. The count and percentage of cells undergoing oxidative stress were quantified using the Muse Cell Analyzer and Muse analysis software (Merck Millipore, USA).

Mitochondrial transmembrane potential detection assay

Mitochondrial transmembrane potential (MMP, ΔΨm) was determined using the MitoPT™ JC-1 assay kit (Immunochemistry Technologies, LCC, MN, USA) according to the manufacturer’s instructions. JC-1 exists as a green fluorescent monomer at low MMP, while at higher MMP, JC-1 forms red fluorescent aggregates and can therefore be
used as a sensitive measure of changes in MMP. Briefly, the cells were resuspended in the MitoPT™ JC-1 solution and incubated in the dark for 15 min at 37 °C. After incubation, the cells were resuspended in 1× assay buffer and seeded in a 96-well clear bottom black plate at a density of 1x10^5 cells/100 μl per well. Fluorescence was assessed in the microplate reader by measuring the JC-1 aggregates (590 nm emission) following 488 nm excitation (JC-1 monomers). The changes in mitochondrial membrane potential were calculated as the ratio of JC-1 aggregate to monomer fluorescence intensity. The mitochondrial membrane potential in the treated cells was expressed as the percentage of that in the control cells (100%).

**Statistical analysis**

The data were expressed as the means ± standard deviation (SD). Statistical analyses were performed using the Student’s t test. All the experiments were repeated at least three times. Results with \( p < 0.05 \) were considered significant.

**Results**

**Effects of HBITC on the cell proliferation**

To determine the potential antiproliferative effect of HBITC, the human SH-SY5Y and U87MG cells were treated with HBITC for 24 and 48 h, and then the BrdU and crystal violet assays were performed. As compared to the cells cultured without HBITC, the cell proliferation of the treated SH-SY5Y cells was inhibited in a dose- and time-dependent manner (Fig. 1a). The inhibition of SH-SY5Y cell proliferation was significant in case of treatment with 40 μM and 60 μM HBITC for 24 h (~ 20% of inhibition) and 48 h (~ 20 and 40% of inhibition, respectively). U87MG cells were less reactive to HBITC—the concentrations up to 60 μM did not affect the proliferation of these cells. At the higher concentration (80 μM), HBITC inhibited U87MG proliferation by ~ 40% after 48 h of culture (Fig. 1b).

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Fig. 1 The effect of HBITC on the proliferation of SH-SY5Y (a) and U87MG (b) cells. SH-SY5Y and U87MG cells were incubated for 24 and 48 h at various concentrations (20, 40, 60, 80 μM) of HBITC. Cell proliferation was analyzed by the BrdU and crystal violet assay.

The data presented in the figure are the mean ± SD of four-five independent experiments. \(* p < 0.05 \) was considered statistically significant as compared with non-treated controls.
Effect of HBITC on the H$_2$S level

Cell imaging experiments were performed to detect H$_2$S in the cells. The cells were incubated with 40 µM and 60 µM HBITC (SH-SY5Y) and with 60 and 80 µM HBITC (U87MG). After incubation with WSP-5 fluorescence probe, strong fluorescence signals were observed (Fig. 2), what confirmed that H$_2$S was released from HBITC.

Effect of HBITC on the thiosulfate level

The results showed that after treating the cells with HBITC for 24 and 48 h, the thiosulfate level in the culture medium was significantly increased (Fig. 3). The thiosulfate level in SH-SY5Y and U87MG cells was about 1.5–2.5-fold and 2-fold of that of the control group, respectively.

Effect of HBITC on the level of proteins involved in cell cycle regulation and H$_2$S metabolism

The Western blot analysis showed that the level of p53, a regulator molecule of cell cycle, was significantly decreased in SH-SY5Y cells upon treatment with HBITC in a dose-dependent manner (Fig. 4a). In addition, the level of p21 was increased after 40 µM HBITC treatment of these cells. Furthermore, the levels of TST and MPST, proteins involved in H$_2$S metabolism, were reduced in SH-SY5Y cells treated with HBITC, but the level of CTH sulfurtransferase was unchanged (Fig. 4a). Interestingly, HBITC did not affect the level of these proteins in U87MG cells (Fig. 4b).

Effect of HBITC on the Bcl-2 expression

A Muse™ Bcl-2 Activation Dual Detection Kit was used to measure the percentage of Bcl-2 protein activation in the cells. The percentage of the cells with active and inactive (phosphorylated) form of Bcl-2, and non-expressing cells was determined. As shown in Fig. 5a–c, the active form...
of Bcl-2 was decreased, but the inactive form of Bcl-2 was increased in both HBITC-treated cancer cell lines (*$p < 0.05$).

**Effect of HBITC on ROS production**

The count and percentage of the cells undergoing oxidative stress were measured by flow cytometry using the Muse Oxidative Stress Kit. We observed that HBITC caused an increase in the number of ROS-positive SH-SY5Y cells (Fig. 6a, b). HBITC did not result in ROS elevation in U87MG cells under the experimental conditions (Fig. 6a, c).

**Effect of HBITC on the mitochondrial transmembrane potential ($\Psi_m$)**

Apoptosis is often accompanied by mitochondrial dysfunction. To investigate the effect of HBITC on the mitochondrial function of SH-SY5Y and U87MG cells, the indicators of mitochondrial activity, such as $\Delta \Psi_m$, were monitored using JC-1 staining. We observed that the incubation of the cells in the presence of HBITC caused a significant decrease in $\Psi_m$ of both cancer cells (Fig. 7).

**Discussion**

Natural isothiocyanates, such as sulphoraphene, benzyl isothiocyanate (BEITC), phenethyl isothiocyanate (PEITC), and iberin, have anti-cancer properties on prostate (Pawlik et al. 2017), breast (Lin et al. 2017), colon (Lai et al. 2010), glioma (Zhu et al. 2013; Su et al. 2015), neuroblastoma (Jadhav et al. 2007), leukemia and lung (Gupta et al. 2014b) cancer cells.

In the present study, it has been demonstrated for the first time that 4-hydroxybenzyl isothiocyanate results in the inhibition of proliferation of human neuroblastoma SH-SY5Y (Fig. 1a) and glioblastoma U87MG (Fig. 1b) cells. Interestingly, we have shown that the anti proliferative effect of HBITC is associated with an increase in $H_2S$ (Fig. 2) and thiosulfate levels (Fig. 3). Hydrogen sulfide exists in the equilibrium with $HS^-$ (in the ratio 2:8) at physiological pH. Furthermore, $H_2S$ (and $HS^-$) has a very short half-life in biological fluids including cell culture medium and blood (Marutani et al. 2015). Thus, it seems that thiosulfate produced from HBITC, and other products of $H_2S$ metabolism, can be involved in the mechanisms that lead to inhibition of cell proliferation.

We have found that HBITC has a more potent inhibitory effect on SH-SY5Y cells proliferation compared to U87MG cells, at the same concentrations in these cells (Fig. 1). Lee et al. (2014) reported that $H_2S$ enhanced glucose uptake and increased glycolytic rate, lactate production, intracellular acidification and cancer cell death (Lee et al. 2014). It appears that U87MG cells, which have a higher than SH-SY5Y cells mitochondrial metabolic capacity, expression of PPAR-$\gamma$ (peroxisome proliferator-activated receptor-$\gamma$) and lactate dehydrogenase (Kim et al. 2015), are to a lesser degree responsible for HBITC-induced changes.

In the present study, the increased $H_2S$ and thiosulfate levels in HBITC-treated SH-SY5Y cells have been associated with downregulation of the level of rhodanese and MPST, but the level of $\gamma$-cystathionase has not been changed in both cancer cell lines (Fig. 4). MPST and rhodanese are localized in the mitochondria and exhibit antioxidant properties (Nagahara 2011; Nagahara et al. 2015). Their catalytic site cysteines are reversibly oxidized by ROS. In the presence of a high level of $H_2S$ and thiosulfate (Figs. 2 and 3), the sulfhydryl groups of proteins can be modified by S-sulfuration. This process is a post-translational modification of l-cysteine residues (-SH to –SSH), by which $H_2S$ exerts various biological effects (Zhang et al. 2017b; Toohey 2012; Mustafa et al. 2009). $H_2S$ induces S-sulfuration of enzymes or receptors, transcription factors, and ion channels.
Fig. 5 Effect of HBITC on the Bcl-2 expression in SH-SY5Y and U87MG cells. The cells were incubated with 20 µM, 40 µM, 60 µM and 80 µM of HBITC for 24 and 48 h. The percentage of the cells with active and inactive form of Bcl-2, and non-expressing cells was determined by flow cytometry analysis using a Muse™ Bcl-2 Activation Dual Detection Kit. a Bar graphs represent the mean ± SD from three-four independent experiments. *p < 0.05 (Student’s t test). b, c Representative results are shown.
Fig. 6 Reactive oxygen species production detected in HBITC-treated SH-SY5Y and U87MG cells. The control cells and the cells treated with HBITC at concentrations of 20, 40, 60, and 80 µM were incubated for 24 and 48 h. The percentage of ROS-positive and ROS-negative cells was determined. a Bar graphs represent the mean ± SD from three independent experiments. b, c Representative results are shown. The blue area (M1) represents the negative cells ROS(−), and the red area (M2) represents the cells with ROS activity [ROS(+)].
disruption of mitochondrial membrane potential and down-regulation of Bcl-2 protein (Fig. 5). It is possible that sulfhydryl groups of Bcl-2 can be modified via S-sulfuration or oxidative stress, as we previously reported (Jurkowska et al. 2017). Interestingly, we have observed in SH-SY5Y cells downregulation of p53 and upregulation of p21 protein levels after HBITC treatment (Fig. 4). Both SH-SY5Y and U87MG cancer cell lines contain wild-type p53 protein (Mustafa Rizvi et al. 2014; Janardhanan et al. 2009; Cerrato et al. 2001; Giacoppo et al. 2017). It seems that inhibition of p53 protein in SH-SY5Y cells can be also associated with S-sulfuration or induction of oxidative stress in these cells. The cysteine residues located at the surface of p53 protein (Cys 124, Cys 176, Cys 182, Cys 229, Cys 242, and Cys 277) function as redox sensors (Kim et al. 2011; Bykov et al. 2016). It was reported that Cys 277 was essential for sequence specific DNA binding and a target for redox regulation of p53 (Buzek et al. 2002). p53 cysteine thiol modification is inversely correlated with its DNA binding properties and transcriptional activity (Kim et al. 2011; Liu et al. 2008). When p53-deficient human lung cancer NCI-H358 cells were transfected with wild-type p53 and subsequently challenged with H2O2, decreased DNA binding properties as well as reduced transcriptional activity of p53 were observed (Parks et al. 1997). Giacoppo et al. (2017) showed that treatment of SH-SY5Y cells with the moringa isothiocyanate complexed with α-cycloextrin (MAC complex) caused inhibition of proliferation and induction of p21 protein, but p53 protein level was decreased at low concentrations of MAC complex. Moreover, Mustafa Rizvi et al. (2014) reported that aluminum-induced apoptosis of SH-SY5Y cells was correlated with increasing ROS production and intracellular calcium level, and with downregulation of p53 protein level.

Our data strongly suggest that upregulation of p21 protein level in SH-SY5Y cells treated with HBITC occurs in a p53-independent manner. In addition to the well-known p53-dependent p21 activation, p21 can be activated by a variety of molecules, i.e. specificity protein 1 (Sp1), histone deacetylase, activating protein-2 (AP-2), elongation 2 factors (E2Fs), signal transducers and activators of transcription (STATs) (Piccolo and Crispi 2012; Roninson 2002). It is known that isothiocyanates are the histone deacetylase inhibitors (Nian et al. 2009). It is, therefore, possible that HBITC treatment results in p21 protein expression level in SH-SY5Y cells, because it affects the histone deacetylase activity.

Conclusions

4-Hydroxybenzyl isothiocyanate, a natural H2S-donor, exerts the antiproliferative effect on SH-SY5Y and U87MG cells. Changes of mitochondrial TST and MPST levels, an increased ROS level, and changes of the level of cell cycle proteins in HBITC-treated SH-SY5Y cells, but not in U87MG cells, are associated with a lower mitochondrial metabolic capacity (Kim et al. 2015) and with a lower level of GSH (Jurkowska et al. 2017) in SH-SY5Y cells. Down-regulation of mitochondrial TST and MPST protein levels...
in the presence of HBITC can result in an increased ROS level in SH-SY5Y cells (Fig. 8). The sulphydryl groups of p53 protein could be modified via HBITC-induced oxidative stress in SH-SY5Y cells, what can trigger a decrease in p53 protein level. It is possible that inactivation of p53 protein in HBITC-treated SH-SY5Y cells can be also caused by its S-sulfuration in the presence of high levels of H2S and thiosulfate. Interestingly, it seems that the induction of p21 protein level in SH-SY5Y cells is associated with p53-inactivation of Bcl-2 protein via HBITC induced membrane potential and number of cells with activated Bcl-2 protein (inactivation of Bcl-2 protein via HBITC induced S-sulfuration or oxidative stress) (Fig. 8).

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Author contributions Conceived and designed the experiments: HJ. Performed the experiments: HJ; DSz; EJ-G. Analyzed the data: HJ. Wrote the paper: HJ, MW.

Compliance with ethical standards

Conflict of interest The authors declare no conflict of interest.

Ethical standard This article does not contain any studies with human participants or animals performed by any of the authors.

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Fig. 8 The antiproliferative activities of 4-hydroxybenzyl isothiocyanate

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