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Cobalt chloride affects the death of SH-SY5Y cells induced by inhibition of ubiquitin proteasome system. Role of heat shock protein 70 and caspase 3

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Abstract. The aim our study was to investigate protective effect of cobalt chloride (CoCl₂) in the model of proteasome stress of neuroblastoma SH-SY5Y cells induced by bortezomib, an inhibitor of 26S proteasome. We have focused our interests on Hsp70 and activation of caspase 3. Finally, we have compared the effect of CoCl₂ with an effect of the pre-treatment of the cells with 17-AAG, an inhibitor of Hsp90 that is capable to induce expression of Hsp70, or with IOX2, an inhibitor of isoform 2 of prolyl hydroxylase that increases stability of hypoxia-inducible factor 1 α (HIF1 α). Pre-treatment of SH-SY5Y cells for 24 h with CoCl₂, at concentrations of 150 or 250 µmol/l, and with 17-AAG at concentration 1 µmol/l but not with IOX2 at concentration 100 µmol/l, was associated with significantly increased expression of Hsp70. We have shown that pre-treatment of SH-SY5Y cells with CoCl₂ but not with 17-AAG or IOX2 was associated with significant delay of the cell death induced by proteasome stress. CoCl₂-mediated effect was consistent with inhibition of bortezomib-induced caspase 3 activation in the cells pre-treated with CoCl₂. Despite established neuroprotective properties of Hsp70 our results do not provide strong evidence that the effect of CoCl₂ could be mainly attributed to the ability of CoCl₂ to induce expression of Hsp70 and other mechanisms have to be considered.

Key words: Neuroprotection — Ubiquitin proteasome system — Cobalt chloride — Heat shock proteins — Caspase 3

Abbreviations: 17-AAG, 17-*N*-allylamino-17-demethoxygeldanamycin; AD, Alzheimer's disease; ALS, amyotrophic lateral sclerosis; CHAPS, 3-[(3-cholamidopropyl)dimethylammonio]-1-propanesulfonate hydrate; HD, Huntington's disease; HIF1α, hypoxia-inducible factor 1α; Hsp, heat shock protein; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; OGD, oxygen glucose deprivation; PD, Parkinson's disease; UPS, ubiquitin proteasome system; VHL, von Hippel-Lindau protein.

Introduction

Ubiquitin proteasome system (UPS) represents an important intracellular pathway involved in the control of a wide range of cellular functions like protein stability, intracellular protein localization, protein-protein interactions, and transcriptional activity (Glickman and Ciechanover 2002). These effects are mediated by monoor poly-ubiquitinylation of specific proteins involved in the regulation of cell cycle, apoptosis, transcription and signal transduction (Schrader et al. 2009). In addition to regulatory functions, the most prominent function of UPS is elimination of aged and aberrant proteins, including misfolded or aggregated proteins. Proteinopathies that include diverse human pathologies including neurodegenerative disorders such as Alzheimer's disease (AD),

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Parkinson's disease (PD), Huntington's disease (HD) and amyotrophic lateral sclerosis (ALS) are characterised by accumulation of certain structurally abnormal proteins and protein aggregates with consequent disruption of cellular functions (Luheshi and Dobson 2009; Chiti and Dobson 2017; Hartl 2017). Proteasome stress resulting from either UPS overload or dysfunction is considered to be important common mechanism implicated in pathophysiology of mentioned neurodegenerative diseases (Ciechanover and Kwon 2015) as well as ischemic neurodegeneration (Luo et al. 2013; Caldeira et al. 2014). The previous research has documented that aggregates of proteins specific for particular neurodegeneration could itself cause strong inhibition of 26S proteasome. The inhibition of proteasome was documented for tau protein associated with AD (Myeku et al. 2016), α -synuclein (Lindersson et al. 2004; Zhang et al. 2008) and parkin (Um et al. 2010) associated with PD, ALS specific proteins (Cheroni et al. 2009) as well as for ubiquitin-conjugated protein aggregates (Bence et al. 2001) produced in early reperfusion phase after global brain ischemia (Racay 2012).

Cobalt in the form of divalent cation is essential to human health since it plays a critical role in the synthesis of vitamin B_{12} . In 1966, the toxic effect of Co^{2+} attributed to the addition of cobalt sulphate to the beer as a stabilizer (Sullivan et al. 1968) was first described among beer drinkers that developed a cardiomyopathy. In addition to heart, Co²⁺ exhibited toxic effects on the cells of other organs including cells of nervous tissue (Caltana et al. 2009; Yang et al. 2011; Guo et al. 2013). The mechanism of the detrimental action of Co^{2+} on the cells is not completely known. Previous studies have demonstrated mitochondrial dysfunction and transmembrane potential collapse caused by opening the mitochondrial transition pore and inhibition of the mitochondrial respiratory chain complexes (Battaglia et al. 2009). In addition, induction of apoptosis (Yang et al. 2004; Jung et al. 2007; Walls et al. 2009; Chang et al. 2016) autophagy (Yang et al. 2015; Fung et al. 2016) and oxidative stress (Chen et al. 2010; Stenger et al. 2011; Guan et al. 2015) has also been documented among the mechanisms of Co^{2+} -induced cell death.

On contrary, pre-treatment of the cells or animals with non-toxic concentrations of $CoCl_2 24$ h prior to lethal insult was documented to be protective in different models of brain ischemia in both neonatal (Bergeron et al. 2000; Jones et al. 2008; Dai et al. 2014) and adult animals (Valsecchi et al. 2011; Wacker et al. 2012) or brain hypoxia injury (Shrivastava et al. 2008). Hypoxia-inducible factor 1 α (HIF1 α) is considered to be essential molecule associated with protective effect of $CoCl_2$ (Sharp et al. 2001; Jones et al. 2013) since $CoCl_2$ is able to stabilize HIF1 α , a key determinant of the cellular response to hypoxia. Stabilisation of HIF1 α is mediated by inhibitory effect of Co^{2+} on the isoforms of prolyl hydroxylase, the enzyme responsible for hydroxylation of HIF1 α protein on specific proline residue. Hydroxylation of HIF1a leads to binding of the von Hippel-Lindau protein (VHL), which recruits an ubiquitin protein-ligase that targets HIF1a for proteasome degradation (Semenza 2011). In addition to stabilisation of HIF1a, Co^{2+} , as the other ions of transition metals, is able to induce expression of heat shock protein 70 (Hsp70) (Koizumi et al. 2013) that also exhibits protective effects in different models of neuronal cell injury (Kelly et al. 2001; Matsumori et al. 2005) and other neurodegenerative conditions (Turturici et al. 2011).

In our previous studies, we have shown that proteasome stress is associated with death of SH-SY5Y cells despite induction of Hsp70 expression that was documented already 4 h after incubation of the cells with bortezomib, inhibitor of 26S proteasome (Klacanova et al. 2016; Pilchova et al. 2017). The aim our study was to investigate impact of the proteasome stress on SH-SY5Y cells pre-treated with CoCl₂ 24 h prior to induction of proteasome stress with bortezomib. We have focused our interests on Hsp70 that was shown to be over-expressed after pre-treatment of the cells with CoCl₂ and activation of caspase 3. Finally, we have compared the effect of CoCl₂ with an effect of pre-treatment of the cells with 17-N-allylamino-17-demethoxygeldanamycin (17-AAG), an inhibitor of Hsp90 that is capable to induce expression of Hsp70, or with IOX2, an inhibitor of isoform 2 of prolyl hydroxylase that increases stability of HIF1a.

Materials and Methods

Sodium dodecylsulphate (SDS), bovine serum albumin (BSA), IOX2, 17-AAG, cobalt chloride and (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) (all Sigma-Aldrich), 3-[(3-cholamidopropyl) dimethylammonio]-1-propanesulfonate hydrate (CHAPS) (ApliChem), bortezomib (Santa Cruz Biotechnology), HALTTM protease inhibitor cocktail (ThermoFisher Scientific), prestained protein standards (BioRad, cat. no. 1610373). Mouse monoclonal antibodies against Hsp27 (SC-13132), Hsp70 (SC-66048), Hsp90 (SC-13119), caspase 3 (SC-271028), HIF1 α (SC-71247) and β -actin (SC-47778) (all Santa Cruz Biotechnology). Goat anti-mouse (A0168) (all Sigma-Aldrich) secondary antibodies conjugated with horse radish peroxidase.

Cell culture and treatment

SH-SY5Y cells (ATCC) were maintained in DMEM:F12 (1:1) medium supplemented with 10% fetal bovine serum, 1% penicillin-streptomycin (all PAA) at an optimal cell density of 0.5×10^6 cells/ml at 37°C and 5% CO₂ humidified atmosphere. The media were changed every 3 days. Naïve

or SH-SY5Y cells pre-treated with CoCl₂, 17-AAG and IOX2 24 h prior to treatment with indicated concentrations of bortezomib (0, 10, 20, 50 nmol/l) for 24 and 48 hours at 37°C and 5% CO₂ humidified atmosphere were at the end of the treatment washed 3 times with ice cold PBS and then re-suspended in a lysis buffer (30 mmol/l Tris-HCl, 150 mmol/l NaCl, 1% CHAPS, 1× protease inhibitor cocktail, pH 7.6) for total protein extraction. Proteat concentrations were determined by protein Dc assay kit (Bio-Rad) using BSA as standard.

Cell viability

Cells were seeded in 96-well plates at concentrations of 0.4×10^6 SH-SY5Y cells *per* ml. Naïve or SH-SY5Y cells pre-treated with CoCl₂, 17-AAG and IOX2 24 h prior to treatment with bortezomib were incubated for 24 and 48 h with indicated concentrations of bortezomib (0, 5, 10, 20, 50 nmol/l) at 37°C in 5% CO₂ humidified atmosphere. At the end of treatment, 0.01 ml of MTT solution (5 mg/ml) were added to each well and the cells were further incubated for 4 hours at 37°C and 5% CO₂ humidified atmosphere. The insoluble formazan, which resulted from oxidation of added MTT by vital cells, was dissolved by addition of 0.1 ml of SDS solution (0.1 g/ml) and overnight incubation at 37°C and 5% CO₂ humidified atmosphere. The absorbance of formazan was determined spectrophotometrically using microplate reader Bio-Rad 2010. The relative viability of the cells was determined as ratio of optical density of formazan produced by treated cells to optical density of formazan produced by non-treated control cells and expressed as percent of control. For each treatment time, the optical density value of nontreated control cells was considered as 100% of viable cells.

Western blotting

Isolated proteins (30 µg of proteins loaded per lane) were separated on 12% SDS-polyacrylamide gels (PAGE) under reducing conditions. Separated proteins were transferred to nitrocellulose membranes using semidry transfer and membranes were probed with antibodies specific to Hsp90 (1:500), Hsp70 (1:500), Hsp27 (1:500), HIF1a (1:200), caspase 3 (1:200) and β -actin (1:1000). Further incubation of membranes with particular secondary antibodies (all 1:5000) was followed by visualization of immunopositive bands using the chemiluminiscent substrate SuperSignal West Pico (Thermo Scientific) and Chemidoc XRS system (Bio-Rad). Intensities of specific bands were quantified by Quantity One software (Bio-Rad). The intensities of bands of interest were normalized to corresponding intensities of bands of β -actin and expressed as intensity of the band of particular protein in treated cells relative to intensity of band in control non-treated cells.

Statistical analysis

For the comparison of differences in protein expression nontreated and pre-treated cells, one-way ANOVA (GraphPad InStat V2.04a, GraphPad Software) was first carried out to test for differences among all experimental groups followed by the Tukey's test to determine differences between individual groups. The value p < 0.05 was considered as being significant.

Results

In order to mimic hypoxic preconditioning, SH-SY5Y cells were pre-treated for 24 h with CoCl₂ at concentrations 150 µmol/l and 250 µmol/l. Incubation of SH-SY5Y cells for 24 h with CoCl₂ at concentrations 150 µmol/l and 250 µmol/l was associated with significantly increased expression of Hsp70 to 829% (*p* < 0.01) and 1321% (*p* < 0.001) of control, respectively (Fig. 1A). Expression of Hsp27 was also significantly elevated to 337% of control (p < 0.01) in the cells incubated for 24 h with CoCl_2 at concentration 250 $\mu mol/l$ while expression of Hsp90 was not significantly altered in the cells incubated for 24 h with CoCl₂ at concentrations 150 µmol/l and 250 µmol/l. The impact of bortezomib on the relative viability of the SH-SY5Y cells and effect of pretreatment of the SH-SY5Y cells with CoCl₂ was assessed by MTT test (Fig. 1B, C). In agreement with our previous results (Klacanova et al. 2016; Pilchova et al. 2017), treatment of SH-SY5Y cells with bortezomib at concentrations 10 nmol/l and higher for 24 and 48 h was associated with decreased relative cell viability (Fig. 1B, C). Pre-treatment of the cells with CoCl₂ 24 h prior to application of bortezomib was associated with significantly higher relative viability of the cells treated 24 h with bortezomib at concentrations 20 nmol/l and 50 nmol/l (Fig. 1B). After 48 h of treatment, the relative viability of the cells pre-treated with CoCl₂ was higher than relative viability of the cells that were not pre-treated however, the difference was not statistically significant (Fig. 1C). Since activation of caspase 3 in SH-SY5Y cells treated with bortezomib for 48 h was clearly documented in our previous study (Pilchova et al. 2017), we have also investigated impact of pre-treatment of SH-SY5Y cells with CoCl₂ on bortezomib-induced activation of caspase 3. As shown on Fig. 1D, activation of caspase 3 in SH-SY5Y cells was documented by decrease of procaspase 3 and appearance of p17 fragment of active caspase 3. These changes were observed 48 h after the treatment of the cells with bortezomib at concentrations 10 and 20 and 50 nmol/l. The level of p17 fragment of active caspase 3 was decreased in the SH-SY5Y cells pre-treated with 150 µmol/l CoCl2 and was not observed in the SH-SY5Y cells pre-treated with 250 µmol/l CoCl2 and then treated with bortezomib at concentrations 10, 20 and

50 nmol/l. In addition, bortezomib-induced decrease of the level of non-active procaspase 3 was almost completely reversed in the SH-SY5Y cells pre-treated with either 150 or $250 \,\mu$ mol/l CoCl₂ and then treated with 10, 20 and 50 nmol/l of bortezomib.

In order to assess the involvement of Hsp70 in the protection of SH-SY5Y cells against proteasome stress we have also used 17-AAG inhibitor of Hsp90 that is known to induce expression of Hsp70 (Powers and Workman 2007). In fact, incubation of SH-SY5Y cells for 24 h with 1 µmol/l 17-AAG was associated with significantly increased expression of Hsp70 to 632% (p < 0.01) of control (Fig. 2A). Expression of Hsp27 was also significantly elevated in the cells incubated with 1 µmol/l 17-AAG to 894% (p < 0.001) of control (Fig.



Figure 1. Impact of CoCl₂ on expression of heat shock proteins, bortezomib-induced cell death and activation of caspase3. A. Total cell extracts were prepared from SH-SY5Y cells after the pre-treatment with indicated concentrations of CoCl2 for 24 h. The effect of CoCl₂ on the levels of Hsp70, Hsp27 and Hsp90 was evaluated by Western blot analysis of total cell extracts as described in Materials and Methods. Data are presented as means \pm SD (n = 4). * p < 0.05, ** p <0.01, *** p < 0.001 (one-way ANOVA, followed by Tukey's test to determine differences at the levels of particular proteins between control non-treated cells and cells pre-treated with CoCl₂). B, C. SH-SY5Y were first pre-treated with indicated concentrations of CoCl₂ and then treated with indicated concentrations of bortezomib for 24 (B) and 48 h (C). After treatment, relative cell viability was determined by MTT test as described in Material and Methods. Data are presented as means ± SEM (4 independent experiments performed in triplicate). * p < 0.05, ** *p* < 0.01, *** *p* < 0.001 (one-way ANOVA followed by Tukey's test to determine differences between relative viability of control cells and cells treated with different concentrations of bortezomib as well as between relative viability of non-pre-treated cells and cells pre-treated with CoCl₂ and then treated with particular concentration of bortezomib). D. Total cell extracts were prepared from SH-SY5Y cells that were either non-pretreated or pre-treated with indicated concentrations of CoCl₂ (150 and 250 µmol/l) and then treated with indicated concentrations of bortezomib 48 h. The activation of caspase 3 was evaluated by Western blot analysis of total cell extracts as described in Materials and Methods.

2A). Expression of Hsp90 was not significantly altered in the cells incubated for 24 h with 17-AAG at concentrations 0.5 µmol/l and 1 µmol/l (Fig. 2A). Despite induction of Hsp70 expression, we did not observe significant differences of relative viability of the cells that were pre-treated with 17-AAG after the incubation with bortezomib in comparison to cells that were not pre-treated with 17-AAG but also incubated with bortezomib (Fig. 2B). In accord with the results of MTT test, pre-treatment of SH-SY5Y cells with 17-AAG did not suppress activation of caspase 3 since p17 fragment of active caspase 3 was observed in the cells treated with bortezomib irrespective of pre-treatment with 17-AAG (Fig. 2C). In addition, decrease of non-active procaspase 3 was observed in both experimental conditions.

Finally, we have pre-treated the cells with IOX2 inhibitor of prolyl hydroxylase 2 that is known to induce expression of HIF1a (Chan et al. 2015). Despite some opposing results, HI-F1a is considered to be an essential molecule associated with neuroprotection in the model of hypoxic preconditioning induced by CoCl₂ (Jones et al. 2013). The level of HIF1a in the cell extracts from control un-treated SH-SY5Y cells was under Western blot detection limit (Fig. 3) but pre-treatment of the cells with IOX2 at concentration 100 µmol/l was associated with stabilization of HIF1a as documented by detection of proper signal on Western blot (Fig. 3A). Pre-treatment of the cells with IOX2 at concentration 100 µmol/l did not significantly alter expression of Hsp70 and Hsp90 (Fig. 3A) but was associated with significantly increased expression of Hsp27 to 186% of control (p < 0.05). Despite stabilization of HIF1a and increased expression of Hsp27, pre-treatment of the SH-SY5Y cells with IOX2 at concentration 100 µmol/l was not associated with significant differences of relative cell viability as documented by MTT test (Fig. 3B).

Discussion

We have shown in this study that pre-treatment of SH-SY5Y cells for 24 h with $CoCl_2$, at concentrations of 150 or 250 µmol/l, and with 17-AAG at concentration 1 µmol/l but not with IOX2 at concentration 100 µmol/l, was associated with significantly increased expression of Hsp70.



Figure 2. Impact of 17-AAG on expression of heat shock proteins, bortezomib-induced cell death and activation of caspase 3. A. Total cell extracts were prepared from SH-SY5Y cells after the pre-treatment with indicated concentrations of 17-AAG for 24 h. The effect of 17-AAG on the levels of Hsp70, Hsp27 and Hsp90 was evaluated by Western blot analysis of total cell extracts as described in Materials and Methods. Data are presented as means ± SD (n = 4). ** p < 0.01, *** p < 0.001 (oneway ANOVA, followed by Tukey's test to determine differences at the levels of particular proteins between control non-treated cells and cells pre-treated with 17-AAG). B. SH-SY5Y were first pre-treated with 17-AAG and then treated with indicated concentrations of bortezomib for 24 h. After treatment, relative cell viability was determined by MTT test as described in Material and Methods. Data are presented as means ± SEM (4 independent experiments performed in

triplicate). * p < 0.05, ** p < 0.01, *** p < 0.001 (one-way ANOVA followed by Tukey's test to determine differences between relative viability of control cells and cells treated with different concentrations of bortezomib as well as between relative viability of non-pre-treated cells and cells pre-treated with 17-AAG and then treated with particular concentration of bortezomib). **C.** Total cell extracts were prepared from SH-SY5Y cells that were either non-pre-treated or pre-treated with 17-AAG and then treated with indicated concentrations of bortezomib 48 h. The activation of caspase 3 was evaluated by Western blot analysis of total cell extracts as described in Materials and Methods.



Figure 3. Impact of IOX2 on expression of heat shock proteins, stabilisation of HIF1a and bortezomib-induced cell death. **A.** Total cell extracts were prepared from SH-SY5Y cells after the pre-treatment with IOX2 for 24 h. The effect of IOX2 on the levels of Hsp70, Hsp27, Hsp90 and HIF1a was evaluated by Western blot analysis of total cell extracts as described in Materials and Methods. Data are

presented as means \pm SD (n = 4). * p < 0.05 (one-way ANOVA, followed by Tukey's test to determine differences at the levels of particular proteins between control non-treated cells and cells pre-treated with IOX2). **B.** SH-SY5Y were first pre-treated with IOX2 and then treated with indicated concentrations of bortezomib for 24 h. After treatment, relative cell viability was determined by MTT test as described in Material and Methods. Data are presented as means \pm SEM (four independent experiments performed in triplicate *per* each cell line). * p < 0.05, ** p < 0.01 (one-way ANOVA followed by Tukey's test to determine differences between relative viability of control cells and cells treated with different concentrations of bortezomib as well as between relative viability of non-pre-treated cells and cells pre-treated with IOX2 and then treated with particular concentration of bortezomib).

Pre-treatment of SH-SY5Y cells with CoCl₂, but not with 17-AAG or IOX2, was associated with significant delay of the cell death induced by proteasome stress. CoCl₂-mediated effect was consistent with inhibition of bortezomib-induced caspase 3 activation in the cells pre-treated with CoCl₂. Despite established neuroprotective properties of Hsp70 our results do not provide strong evidence that the effect of CoCl₂ could be mainly attributed to the ability of CoCl₂ to induce expression of Hsp70 since pre-treatment of the cells with 17-AGG.

Cobalt in the form of divalent cation is an essential microelement that at higher concentrations exhibits significant toxicity to the cells of different tissues including cells of nervous system. Detrimental effects of the Co²⁺ were among other mechanisms attributed to the induction of mitochondrial apoptosis (Lee et al. 2008) including activation of caspase 3 (Zou et al. 2002; Peng et al. 2015; Sun et al. 2015). On contrary, our results have shown that delay of bortezomib-induced cell death might be a result of inhibition of bortezomib-induced caspase 3 activation that was documented in the SH-SY5Y cells pre-treated with subtoxic concentrations of CoCl₂. Delay of the cell death after inhibition of caspases was documented in several previous studies. For example, fast apoptotic death of primary hippocampal and cortical neurones induced by ABT-737 that was shown to induce Bax/Bak-dependent mitochondrial apoptosis (Vogler et al. 2009) was not prevented but was significantly delayed by application of the broad spectrum caspase inhibitor zVADfmk (Young et al. 2010). In accord with our results, CoCl₂ at concentrations 100 and 200 µmol/l blocked isocudraxanthone-induced apoptosis and activation of caspase 3 (Shin et al. 2014). Concentration of $CoCl_2$ seems to be critical with respect to the impact of $CoCl_2$ on the exposed cells. Pre-treatment of neuronal cells with $CoCl_2$ at concentration 100 µmol/l for 20 h prior to exposure of the cells to oxygen glucose deprivation (OGD) attenuated OGD-induced neuronal death whereas treatment of the cells with $CoCl_2$ at concentrations 300 or 500 µmol/l induced significant neuronal death, even without exposure to OGD (Jones et al. 2013). Similarly, activation of caspase 3 was documented in mouse cortical HT-22 neurones treated with $CoCl_2$ at concentrations 300 µmol/l and higher (Peng et al. 2015).

It was shown in numerous of the previous studies that heat shock proteins can suppress protein aggregation (Doyle et al. 2013) and toxicity in different models of neurodegenerative diseases (Stetler et al. 2010; Lindberg et al. 2015; Smith et al. 2015). With respect to apoptosis initiation, Hsp70 was shown to act at the level of extrinsic apoptosis (Clemons et al. 2005) as well as caspase-dependent (Beere et al. 2000; Saleh et al. 2000; Stankiewicz et al. 2005) or caspase-independent (Ravagnan et al. 2001) intrinsic (mitochondrial) apoptosis. With respect to neuronal cells, Hsp70 over-expression in rat primary cortical neurons and the SH-SY5Y cells protected the cells in four independent models of apoptosis: etoposide-, staurosporine-, C2-ceramide-, and β-amyloid-induced apoptosis (Sabirzhanov et al. 2012). It was demonstrated that Hsp70 binds and potentially inactivates apoptotic proteaseactivating factor 1 as well as apoptosis-inducing factor, key molecules involved in the development of caspase-dependent and caspase-independent cell death, respectively (Sabirzhanov et al. 2012). Over-expression of Hsp70 in dopaminergic

neurones of *Drosophila* protected them from paraquatinduced cell death *via* inhibition of caspase 3 activation (Shukla et al. 2014). In addition, Hsp70 is considered to be key molecule conferring protection of CA1 pyramidal cells after preconditioned global brain ischemia (Burda et al. 2003; Tanaka et al. 2004) that is also associated with massive proteasome stress (Racay 2012). In our previous study, we have shown that the sensitivity to bortezomib and kinetics of death of leukemic cells correlated well with higher expression of Hsps in resistant and slowly responding K562 cells (Kliková et al. 2015). Thus, CoCl₂-induced massive expression of Hsp70 might represent plausible molecular mechanism responsible for inhibition of caspase 3 activation and delay of bortezomibinduced death of SH-SY5Y cells.

To prove the involvement of Hsp70 in protection from proteasome stress, we have also pre-treated SH-SY5Y cells with 17-AAG, inhibitor of Hsp90 that was shown to induce of Hsp70 expression (Powers and Workman 2007) and to protect effectively the cells from different form of cellular stress including proteasome stress caused by proteasome inhibition (Bonner et al. 2010). In addition, 17-AAG was tested as neuroprotective molecule in several models of neurodegeneration and pre-clinical studies of neurodegenerative diseases. In our experiments, we did not observe protective effect of 17-AAG. The inability of 17-AAG to protect SH-SY5Y cells from proteasome stress observed in our experiments might be explained by lower extent of Hsp70 induction after pretreatment of the cells with 17-AAG as compared to the levels of Hsp70 induced by CoCl₂. On contrary, despite induction of protective Hsp70 17-AAG exhibits also cytotoxic effects and was tested for treatment of different cancers (Chatterjee and Burns 2017). Although, we did not incubate cells simultaneously with 17-AAG and bortezomib, it is important to note that combination of 17-AAG with bortezomib was associated with synergistic cell death effect on U266 (Duus et al. 2006) and MCF-7 (Mimnaugh et al. 2004) cell lines.

Our results open the possibility that there are some other factors beyond Hsp70 that contributed to the protective effect of CoCl₂. For example, Hsp27 was considered to be more protective than Hsp70 in the model of a-synucleininduced neuronal cell death (Zourlidou et al. 2004). Hsp27 seems to exert its protective functions via inhibition of extrinsic mitochondria-independent pathway (Tan et al. 2009). Interestingly, the same study showed the protective effect of Hsp27 against CoCl₂-induced apoptosis (Tan et al. 2009) that depends predominately on initiation of mitochondrial apoptosis. In our experiments, expression of Hsp27 was significantly increased after pre-treatment of cells with 250 µmol/l of CoCl₂ but not with 150 µmol/l CoCl₂. In addition, Hsp27 was shown to be over-expressed in the cells pre-treated with IOX2. Since pre-treatment of the cells with IOX2 was not associated with protective effect and pretreatment with 150 µmol/l of CoCl2 was associated with very similar results as pre-treatment with 250 μ mol/l of CoCl₂, we consider the involvement of Hsp27 in the effects associated with CoCl₂ pre-treatment as unlikely.

Another possibility how to explain effect of CoCl₂ consists in the ability of CoCl₂ to inhibit prolyl hydroxylases that is associated with stabilisation HIF1a (Jones et al. 2013). To investigate possible involvement of HIF1a we have pre-treated cells with IOX2 inhibitor of prolyl hydroxylase 2. The ability of IOX2 to stabilize HIF1a depends on cell type (Chan et al. 2015) but we have clearly documented stabilization of HIF1a after pre-treatment the SH-SY5Y cells with IOX2 at concentration 100 µmol/l. HIF1a is considered to be critical molecule responsible for protective effect of CoCl₂induced hypoxic preconditioning in different models of ischemic brain injury (Sharp et al. 2001; Jones et al. 2013). On contrary, previous studies indicated that CoCl₂ activates HIF1a, acting as a hypoxia-mimetic and inducing reactive oxygen species-mediated toxicity (Chen et al. 2010; Guan et al. 2015). Despite clear stabilisation of HIF1a level after pre-incubation of the cells with IOX2 at concentration 100 µmol/l we have observed only marginal non-significant effect of this pre-treatment. Thus involvement of HIF1a alone in neuroprotective mechanisms associated with CoCl₂ pre-treatment seems to be unlikely but the combined protective effect of both Hsp70 and HIF1a cannot be completely excluded. Finally, other mechanisms activated by CoCl₂, e.g. modulation of expression of Bcl2 family proteins (Shin et al. 2014), could to contribute to protective effects of CoCl₂.

In conclusion, we have shown in this study that pretreatment of SH-SY5Y cells with $CoCl_2$ was associated with significant delay of the cells death induced by proteasome stress. In addition, we have shown that $CoCl_2$ -mediated protective effect was most probably a result of inhibition of bortezomib-induced caspase 3 activation but the result presented in our study cannot be exclusively attributed to $CoCl_2$ -dependent induction of Hsp70.

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