HOMOCYSTEINE AFFECTS CARDIOMYOCYTE VIABILITY: CONCENTRATION-DEPENDENT EFFECTS ON REVERSIBLE FLIP-FLOP, APOPTOSIS AND NECROSIS

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Homocysteine affects cardiomyocyte viability: concentration-dependent effects on reversible flip-flop, apoptosis and necrosis

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Published online: 18 April 2007
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Abstract

Background Hyperhomocysteinaemia (HHC) is thought to be a risk factor for cardiovascular disease including heart failure. While numerous studies have analyzed the role of homocysteine (Hcy) in the vasculature, only a few studies investigated the role of Hcy in the heart. Therefore we have analyzed the effects of Hcy on isolated cardiomyocytes.

Methods H9c2 cells (rat cardiomyoblast cells) and adult rat cardiomyocytes were incubated with Hcy and were analyzed for cell viability. Furthermore, we determined the effects of Hcy on intracellular mediators related to cell viability in cardiomyocytes, namely NOX2, reactive oxygen species (ROS), mitochondrial membrane potential (ΔΨm) and ATP concentrations.

Results We found that incubation of H9c2 cells with 0.1 mM D.L-Hcy (= 60 μM L-Hcy) resulted in an increase of ΔΨm as well as ATP concentrations. 1.1 mM N,N-Hcy (= 460 μM L-Hcy) induced reversible flip-flop of the plasma membrane phospholipids, but not apoptosis. Incubation with 2.73 mM N,N-Hcy (= 1.18 mM L-Hcy) induced apoptosis and necrosis. This loss of cell viability was accompanied by a thread-to-grain transition of the mitochondrial reticulum, ATP depletion and nuclear NOX2 expression coinciding with ROS production as evident from the presence of nitrotyrosine residues. Notably, only at this concentration we found a significant increase in S-adenosylhomocysteine which is considered the primary culprit in HHC.
Conclusion  We found concentration-dependent effects of Hcy in cardiomyocytes, varying from induction of reversible flip-flop of the plasma membrane phospholipids, to apoptosis and necrosis.

Keywords  Hyperhomocysteinaemia · Cardiomyocytes · Heart failure · Oxidant stress · Apoptosis

Introduction

Hyperhomocysteinaemia (HHC) reflects an impairment of the methionine-homocysteine metabolism. Normally homocysteine (Hcy) is converted to cysteine via transulfuration or to methionine by addition of a methyl group donated by folate [1]. However, when these pathways are deranged because of genetic mutations of enzymes involved or relative deficiencies of folate, vitamin B6 or vitamin B12, the serum concentration of Hcy increases [2, 3].

In 1969, McCully et al. reported elevated concentrations of Hcy in young patients with severe cardiovascular disease [4]. They hypothesized the existence of a pathogenic link between HHC and atherogenesis. Since then, other studies have also suggested that HHC is an independent risk factor for cardiovascular complications [5–8]. It is now assumed that even mild HHC constitutes an independent risk factor for cardiovascular disease [9] with equal predictive strength as hypercholesterolemia and smoking [10]. Whether Hcy itself or a Hcy-related substance such as S-adenosyl-homocysteine (SAH) is the culprit, and what the exact pathogenic mechanism of the cardiovascular complications induced by HHC could be, remains to be elucidated. In addition, large-scale intervention trails using high-dose-B-vitamins to reduce plasma Hcy levels in cardiovascular patients have as yet not shown an overall clinical benefit [11–13]. This raises further questions about the mechanisms underlying the association between HHC and cardiovascular disease, and about the possible adverse effects of high-dose-B-vitamins as a means of reducing Hcy levels [14].

Studies of the effects of Hcy on the vascular wall, especially on endothelial and smooth muscle cells, indicate that increased Hcy leads to cellular stress and also to cell damage [15–17]. Hcy primes human neutrophils for an increased production of reactive oxygen species (ROS) and induces ROS production in endothelial cells [15, 18]. ROS production has also been found in the vascular wall of coronary arteries during HHC [19] and a role of NADPH oxidases in this process has been suggested.

From the Framingham Heart Study it is known that increased plasma homocysteine concentrations, next to their effect on the vasculature, also form an independent risk factor for the development of heart failure [20]. Several other studies have also shown a link between HHC and heart failure [21, 22]. However, so far no studies have addressed possible effects of Hcy on cardiomyocytes. Therefore, we performed the present study to analyze the effects of Hcy on isolated cardiomyocytes.

Materials and methods

Cell cultures

Rat cardiomyoblasts (H9c2 (2–1) cells; ATCC, Manassas, VA, USA) were cultured in Dulbecco’s Modified Eagles Medium (DMEM; BioWhittaker, Verviers, Belgium) supplemented with 10% heat-inactivated fetal calf serum (FCS; BioWhittaker), 2 mM l-glutamine (GIBCO, Paisley, UK), 100 IU/ml penicillin (Yamanouchi Europe BV, Leiderdorp, Netherlands) and 100 μg/ml streptomycin (Radiopharma-Fisiopharma, Palomonte, Italy), and grown at 37°C in a humidified 5% CO₂/95% air atmosphere. Experiments were performed with cells grown to a confluency of 60–80%. Adult rat cardiomyocytes were isolated as described before [23].

Antibodies and chemicals

Monoclonal Antibody 48 against NOX2 IgG 1 (dilution 1:28) was obtained from Sanquin Research at CLB, Amsterdam, The Netherlands [24]. Cy3-labeled goat-anti-rabbit-Ig (1:75) (Alexa Fluor; Leiden, Netherlands) was used as a secondary antibody. Polyclonal z-Nitrotyrosine IgG (1:50) (Molecular Probes Inc, Eugene, OR, USA) was used as an antibody to measure nitrotyrosin residues which is an indicator for ROS production. In this case, Cy5-labeled goat-anti-rabbit-IgG (1:50) (Jackson Immuno Research, West Grove, PA, USA) was used as a secondary antibody. Isotype controls and PBS were used to determine aspecific binding. Monoclonal antibody 150/I against pan-actin (1:600; Chemicon International, Inc, Temecula, CA, USA) was used as a loading control for the western blot analysis.

Cells were incubated with several concentrations (0.1 mM–2.7 mM) of reduced homocysteine (Hcy) (Sigma, St. Louis, MO, USA) in growth medium for 24 h at 37°C in a humidified 5% CO₂/95% air atmosphere.

Measuring Hcy concentration in growth medium

The H9c2 cells were incubated with different concentrations of reduced Hcy. Several studies have suggested that only the l form is bioactive [25, 26]. Next to d,l-Hcy concentration we therefore also measured the l-form. d,l-Hcy was measured by HPLC using fluorescence detection [27].

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Intra- and interassay coefficients of variation (CVs; std dev/average *100%) were 1.8 and 3.5%, respectively.
t-Hcy was measured by the Abbott IMx fluorescence polarization immunoassay (IMx; Abbott Laboratories, Abbott Park, IL, USA). Intra- and interassay CVs were less than 2 and 4%, respectively. The concentration of Hcy in culture medium was measured before incubation (t = 0) and after 24 h (final timepoint for the rest of the experiments of incubation with H9c2 cells to determine the amount of putative degradation and/or uptake of the Hcy during this incubation period.

Determination of intracellular S-Adenosylmethionine (SAM) and S-Adenosylhomocysteine (SAH)

Since previous studies suggested that an increase in SAH is the culprit in cellular damage due to elevated levels of Hcy, we also determined the intracellular concentration of SAM and SAH in H9c2 cells after 24 h of incubation with t-Hcy. Tandem mass spectrometry (MS/MS) was used for the determination of SAM/SAH concentrations as previously described [28].

Flow cytometry

Phosphatidylserine exposure was assessed with FITC-labeled human recombinant annexin-V (Bender Med Systems, Vienna, Austria), while propidium iodide (PI; Bender Med Systems) was used to determine the permeability of the cellular membrane. Experiments were performed in the presence or absence of Z-Asp-Glu-Val-Asp-fluoromethylketone (Z-VAD FMK, 25 μM, Alexis Biochemicals, Lausen, Switzerland), a general caspase inhibitor, to differentiate between reversible flip-flop of plasma membrane phospholipids, apoptosis, and necrosis [23]. After treatment with Hcy, the cells were trypsinized and centrifuged at 400g for 5 min at room temperature. Cells were then washed with serum-free DMEM, and resuspended in serum free DMEM containing annexin V (1:40) for 30 min in the dark at 37°C in a humidified 5% CO2/95% air atmosphere. Cells were subsequently washed and resuspended in serum free DMEM containing PI (1:40). Cells were measured with a FACS Calibur (Becton Dickinson, San Jose, CA, USA). Results were analyzed by Cell Quest Pro software (Becton Dickinson).

Detection of caspase-3 activity

Cells were grown in a 96-wells plate (20,000 cells/well). After treatment with Hcy and/or Z-VAD FMK, cells were lysed and incubated with DEVD-Rhodamine 110 substrate (Roche, Mannheim, Germany) for 1 h at 37°C. Subsequently the amount of free rhodamine was determined at a microplate fluorescence reader (TECAN spectralflo, Switzerland). The developed fluorochrome was proportional to the concentration of activated caspase-3 and could be quantified by a calibration curve of diluted free rhodamine. Each condition was measured in triplo per measurement (total of three measurements).

Immunofluorescence microscopy

To measure the expression of NOX2 and the putative formation of nitrotyrosin, cells were incubated with or without Hcy for 24 h in the 4-well chamber slides (Nalgene Nunc International, Naperville, IL, USA). Cells were washed with PBS and fixed with 4% formaldehyde for 10 min at 37°C. Cells were subsequently washed with PBS, permeabilized with acetone-methanol (70%-30%) for 10 min at -20°C, and then washed again with PBS/Tween-20 (0.05% v/v) Tween-20 in PBS). Subsequently cells were incubated with primary antibodies against NOX2 and nitrotyrosin for 60 min at room temperature followed by incubation overnight at 4°C. PBS and isotype controls were also tested to determine nonspecific binding of the antibodies and background signal. The following day the wells were washed with PBS/Tween and incubated with the secondary antibodies for 30 min at room temperature. After subsequent washes in PBS/Tween and PBS, the slides were covered with mounting medium containing DAPI (Vector Laboratories Inc, Burlingame, CA, USA) to visualize nuclei. Thereafter the slides were covered with coverslips.

Subsequently, cells were analyzed by means of a 3i Marianas™ digital imaging microscopy workstation (Zeiss Axiovert 200 M inverted microscope; Carl Zeiss, Sliedrecht, Netherlands), equipped with a nanostepper motor (Z-axis increments; 10 nm) and a cooled CCD camera (Cooke Semicam, 1280 x 1024 pixels; Cooke Co, Towanda, NY, USA). Visualization of NOX2 and nitrotyrosin was performed with a 40x air lens. The microscope, camera and data viewing process were controlled by SlideBook™ software (version 4.0.8.1; Intelligent Imaging Innovations, Denver, CO, USA).

Live cell analysis of Δψm

To visualize mitochondrial membrane potential (Δψm) via Life Cell Imaging, cells were loaded for 30 min at 37°C with 0.5 μmol/L of 5,5',6,6'-tetrachloro-1',3',3'-tetraethylbenzimidazolylcarbocyanine iodide (JC-1; Cell Technology Inc, Minneapolis, USA). In non-apoptotic cells, JC-1 exists as a monomer and is visible on the green channel (FITC). In healthy cells these monomers accumulate
as aggregates in the mitochondria due to existing ΔΨm. These aggregates are visible on the red channel (CY3) when viewed with a fluorescence microscope. Cells were washed with PBS before viewing and subsequently analyzed by use of a Z1 Marians™ digital imaging microscopy workstation as described above.

Western blot analysis

After treatment H9c2 cells were harvested into modified ELB lysis buffer (250 mM NaCl, 0.1% Nonidet P-40, 50 mM HEPES pH 7.0, 5 mM EDTA, 0.5 mM DTT) wherein protease inhibitor cocktail (PIC, 1:40; Sigma) was added and the cell suspension was mixed rigorously and then incubated for 30 min on ice. After determination of the protein concentration of the samples with the BCA protein assay kit (Pierce, Rockford, IL, USA), reducing sample loading buffer (0.25 M TRIS pH 6.8, sodium dodecyl sulfate (SDS), glycerol, 2-mercaptoethanol, bromophenol blue) was added and the samples were mixed and heated at 95°C for 10 min. A total of 30 μg protein of each sample was then subjected to SDS-PAGE, transferred to nitrocellulose membranes and analyzed for NOX2 expression with monoclonal antibody 48 (1:250 dilution) and for pan-actin with monoclonal antibody 1501R as a loading control (1:6000), followed by horseradish-peroxidase-conjugated rabbit-anti-mouse immunoglobulins (RZM-HRP; 1:1000 dilution; DakoCytonation, Glostrup, Denmark). The blots were visualized by enhanced chemo-luminescence (ECL; Amersham Biosciences AB; Uppsala, Sweden). Staining was quantified with a charge-coupled device camera (Fuji Science Imaging Systems; Düsseldorf, Germany) in combination with AIDA Image Analyzer software (Isotopenmessgerät; Staubenhardt, Germany).

ATP measurement via Luciferase–Luciferin assay

After treatment cells were collected by trypsinization and centrifugation at 400g for 5 min. Cells were then counted and equal amounts were taken per condition. After centrifugation for 2 min (400g) the supernatant was discarded. The pellet was resuspended in 150 μL of ice-cold perchloric acid (PCA) 0.4 M. Cells were left on ice for 30 min and then centrifuged again for 5 min (2000g) at 4°C. To the isolated supernatant 7.5 μL of K₂CO₃ 5 M was added to neutralize pH. Samples were subsequently stored at −80°C until measuring. The amount of ATP in the samples was determined with a Luciferase–Luciferin assay (Biafinit GmbH & Co KG, Kassel, Germany) according to the manufacturers’ protocol. Samples were measured in a FluoroNunc MaxiSorp plate (Nalgene Nunc International, Rochester, NY). Luminescence was measured by using a Tecan GENios Plus reader (Tecan Benelux, Mechelen, Belgium).

Statistics

Statistics were performed with the SPSS statistics program (windows version 9.0). To evaluate whether observed differences were significant, One Way ANOVA with post hoc Bonferroni tests were used. All values are expressed as mean ± standard error of the mean (SEM). A P value (two sided) of 0.05 or less was considered to be significant.

Results

Measurement of the concentration of Ω1-homocysteine (Ω1-Hcy), Ω2-homocysteine (Ω2-Hcy), S-adenosyl methionine (SAM) and S-adenosyl homocysteine (SAH)

We tested the effects of Ω2-Hcy at concentrations of 0.1 mM, 1.1 mM and 2.7 mM, in accordance with previous studies in isolated vascular cells [15, 17, 26, 29–31]. We quantified the exact concentration of Ω2-Hcy in medium added to the cells, via HPLC. As depicted in Table 1, the concentration of Ω2-Hcy was according to the concentration added to the growth medium.

Since previous studies have shown that only the Ω form of Hcy is bioactive [25, 26], we also determined the Ω-Hcy concentration in medium by Ixs. We found that 42.7%, 42.5% and 43.3% of the total Ω1-Hcy amount 0.1 mM, 1.1 mM and 2.7 mM, respectively, in fact was Ω-Hcy at t = 0. Furthermore, we determined the concentrations of Ω-Hcy at 24 h, the final time point of incubation, and found a significant decrease in the concentrations of Ω-Hcy, namely for 1.1 mM a significant decrease of 0.14 mM Ω-Hcy; *P < 0.001) and for 2.7 mM a significant decrease of 0.34 mM Ω-Hcy; *P = 0.0301 (P < 0.001).

It has been well documented that increased levels of SAH inhibit several important methylation reactions [32–34]. Therefore we also determined the intracellular concentrations of both SAH and SAM, being one of the primary methyl donors (Table 1). Measuring intracellular concentrations of SAM and SAH in H9c2 cells incubated with and without Ω2-Hcy showed a significant depletion of SAM at 2.7 mM Ω2-Hcy compared to untreated cells and 0.1 mM Ω2-Hcy treated cells (P = 0.006; n = 3). While the concentration of SAH increased significantly at this same concentration compared to all other conditions (P < 0.001; n = 3). Therefore we also examined the effects of the higher, non-physiological concentrations of Hcy.
Table I: Concentration of ν, ω-Hcy and 1-Hcy in culture medium on t = 0, 1-Hcy after 24 h incubation, and intra-cellular concentration of SAM/SAH after 24 h incubation

<table>
<thead>
<tr>
<th>mM Hcy added</th>
<th>mM Hcy measured</th>
<th>[SAM] (μM) t = 24</th>
<th>[SAH] (μM) t = 24</th>
</tr>
</thead>
<tbody>
<tr>
<td>ν, ω-Hcy</td>
<td>[ω-Hcy] t = 0</td>
<td>0.0007 (+/−0.0009)</td>
<td>66.4 (+/−35.6)</td>
</tr>
<tr>
<td>0.1</td>
<td>0.06 (+/−0.01)</td>
<td>0.06 (+/−0.01)</td>
<td>18.9 (+/−6.8)</td>
</tr>
<tr>
<td>1.1</td>
<td>0.46 (+/−0.03)</td>
<td>31.6 (+/−5.2)</td>
<td>16.9 (+/−6.8)</td>
</tr>
<tr>
<td>2.7</td>
<td>1.18 (+/−0.05)</td>
<td>20.4 (+/−7.5)</td>
<td>16.8 (+/−3.4)</td>
</tr>
</tbody>
</table>

All mediators were measured in n = 3. A significant decrease of 1-Homocysteine (1-Hcy) can be seen at 1.1 mM and 2.7 mM after 24 h (* P < 0.001). At 2.7 mM a significant decrease of S-adenosylmethionine (SAM, ** P = 0.006) was detected, compared to all other concentrations (SAH, *** P < 0.001). The supplementation of 1-Hcy at 2.7 mM was only present at 0.1 mM 1, 3, 5-Hcy.

Effect of Hcy on cell viability

We analyzed the effect of Hcy on H9c2 cell viability by measuring annexin V and/or PI positivity using flow-cytometry. Incubation with 0.1 mM 1, 3, 5-Hcy had no effect on single-annexin-V-positivity (Fig. 1A); double-annexin-V, PI positivity (Fig. 1B) or single-PI positivity (not shown, approximately 5% for each condition). In contrast, incubation with 1.1 mM 1, 3, 5-Hcy resulted in a significant increase in single-annexin-V-positive cells (P < 0.002) (Fig. 1A). This increase in single-annexin-V-positive cells was only partly, but not significantly inhibited by Z-VAD FMK (31.6% decrease). 1.1 mM 1, 3, 5-Hcy had no significant effect on the percentage of double-annexin-V/PI-positive cells (Fig. 1B) or on the percentage of single-PI-positive cells (not shown). Incubation with 2.7 mM 1, 3, 5-Hcy, however, resulted in a significant increase of single-annexin-V positive cells (P < 0.001) and double-annexin-V/PI-positive cells (P < 0.044) that could not be inhibited significantly by Z-VAD FMK. Again, we did not see any significant increase in single-PI-positive cells (not shown).

Thus, depending on the concentration of Hcy, either reversibly flip-flopped cardiomyocytes and/or late apoptotic/necrotic cells can be found.

Caspase-3 activity

To determine whether the population of double-annexin-V/PI-positive cells consists of late apoptotic cells or necrotic cells, we performed an assay of caspase-3 activity. Incubation with 0.1 mM and 1.1 mM 1, 3, 5-Hcy did result in a small, but not significant increase in caspase-3 activity that was completely inhibited when Z-VAD FMK was added. Incubation with 2.7 mM 1, 3, 5-Hcy for 24 h, however, induced a significant increase of caspase-3 activity (P < 0.001) that was also completely inhibited with Z-VAD FMK (Fig. 2). Caspase-3 activity measured at earlier time points (6 and 16 h) of Hcy incubation was not significantly increased (results not shown). Thus, significant apoptosis induction of H9c2 cells was only present at 2.7 mM 1, 3, 5-Hcy.
Effect of Hcy on NOX2 expression and nitrosylation

In a previous study we have found that ischemia of cardiomyocytes induced nuclear NOX2 expression coinciding with nuclear nitrosylsine residues, resulting in subsequent apoptosis of these cells [35]. Given the pro-apoptotic effect of 2.7 mM L-Hcy, we analyzed by Western blotting the effect of Hcy on NOX2 expression. We found that NOX2 was constitutively present in whole cell lysates of H9c2 cells (Fig. 3). NOX2 expression was slightly increased at 0.1 mM L-Hcy and increased even more at concentrations of 1.1 mM and 2.7 mM of L-Hcy. It is noteworthy that although we found this increase in NOX2 expression, we were not able to detect apoptosis at the lower concentrations of Hcy. Therefore we subsequently analyzed the intracellular localization of NOX2 with digital imaging. In the cytosol, we found diffuse positivity for NOX2 at all concentrations of Hcy (data not shown). Incubation with 0.1 mM and 1.1 mM, but especially with 2.7 mM L-Hcy resulted in a significant increase in NOX2 expression in the nucleus ($P < 0.001$; $P = 0.047$; $P < 0.001$, respectively) (Fig. 4A). These results appear in contrast with the caspase-3 measurements. We therefore also analyzed the presence of nitrosylsine residues as an indicator for ROS production. We found that neither 0.1 mM nor 1.1 mM L-Hcy resulted in significant nitrosylsine formation in the nucleus. In contrast, 2.7 mM L-Hcy did induce a significant formation of nitrosylsine residues in the nucleus ($P < 0.001$) (Fig. 4B). These results underline that not only nuclear NOX2 expression but also nuclear ROS production is necessary for the induction of apoptosis. It has to be noticed that especially at 2.7 mM L-Hcy a significant loss of approximately 50% of the cells was found ($P < 0.001$, results not shown). As these cells theoretically are predominantly double-annexin-V/PI or single-PI-positive, we conclude that the effect of incubation with 2.7 mM L-Hcy on the amount of double-annexin-V/PI or single-PI-positive cells (see Fig. 1) in all probability is underestimated.

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the nucleus \((P < 0.022; \text{Fig. 5D})\) but no significant nitrotyrosine formation. Incubation with 0.1 mM D,L-Hcy led to significant nuclear NOX2 expression and presence of nitrotyrosine residues \((P < 0.01; \text{Fig. 5A–E})\). As with H9c2 cells, we found only little NOX2 positivity in the cytosol under all Hcy concentrations analyzed (arrow in Fig. 5A).

ATP depletion

An important determinant in the process of cell death is ATP, as it is for instance necessary to complete the process of apoptosis. We therefore measured ATP concentrations in H9c2 cells (Fig. 6). Incubation with 0.1 mM D,L-Hcy resulted in a significant increase in ATP concentration \((P = 0.05)\). In contrast, incubation with 1.1 mM D,L-Hcy led to a decrease in ATP concentration that was not significant compared to control cells, but was significant compared to ATP concentration found in cells that were incubated with 0.1 mM D,L-Hcy \((P = 0.024)\). Incubation with 2.7 mM D,L-Hcy induced an more pronounced decrease of ATP concentration that was significant also when compared to control cells \((P = 0.036)\).

\[ \Delta \Psi_m \]

To explain the contrasting effects of different Hcy concentrations on ATP concentrations, we analyzed the mitochondrial membrane potential \((\Delta \Psi_m)\) with JC-1 staining in H9c2 cells. We found an increase in red signal \((P = 0.006)\) when the H9c2 cells were incubated with 0.1 mM D,L-Hcy compared to control cells (Fig. 7A/B) which is indicative for more actively respiring mitochondria. \(\Delta \Psi_m\) after incubation with 1.1 mM and 2.7 mM D,L-Hcy did not differ significantly from those in control cells (Fig. 7A–E). A difference in the morphology of mitochondria was also visible. In control cells a heterozygous mitochondria population is visible consisting of round and elongated mitochondria. Incubation with 0.1 mM D,L-Hcy showed mostly elongated mitochondria, whereas incubation with 1.1 mM and 2.7 mM D,L-Hcy resulted in a more rounded morphology of the mitochondria, the so called thread-to-grain transition which is seen as an early marker for apoptosis [36].

**Discussion**

In the present study we examined the effects of Hcy on cell viability of cardiomyocytes and found different effects, depending on Hcy concentration (Fig. 8).

The lower concentration of 0.1 mM Hcy resulted in mitochondrial hyperpolarization, an increase of ATP and translocation of NOX2 to the nucleus, but without local nitrotyrosin formation. Mitochondrial hyperpolarization and an increase in ATP concentration have been described as first signs of mitochondrial disturbance on the pathway to apoptosis [36, 37]. On the other hand, they have also been described as part of a cell-protective mechanism [38]. Therefore, the effect of this lower Hcy concentration may be a cellular state characterized by early activity of both pro- and anti-apoptotic pathways, which does not necessarily result in cell death.

In contrast, the intermediate concentration of 1.1 mM Hcy led to rounding of mitochondria, a slight, non significant decrease of \(\Delta \Psi_m\) with a visible thread-to-grain transition of the mitochondrial reticulum, a slight decrease in ATP concentrations, a translocation of NOX2 to the nucleus, again without presence of nitrotyrosin residues, and a flip-flop of the cell membrane phospholipids as
**Fig. 5** Effect of Hcy on intracellular localization of NOX2 and ROS production in adult rat cardiomyocytes. Digital Imaging microscopy pictures of adult rat cardiomyocytes incubated with 0.1 mM t,t,-Hcy during 24 h. Red color (A) indicates NOX2 expression which is visible in the nucleus and in the cytosol (arrow). Green color (B) represents nitrotyrosine formation. The merge image (C) also shows blue DAPI staining for DNA which shows co-localization of NOX2 expression and nitrotyrosine formation in the nucleus. (D) Amount of nuclear NOX2 expression in adult rat cardiomyocytes in control, 0.07 mM and 0.1 mM t,t,-Hcy samples (n = 3). (E) Amount of nitrotyrosine formation in adult rat cardiomyocytes in control, 0.07 mM and 0.1 mM t,t,-Hcy sample (n = 3). The arbitrary units are defined as sum intensity per nucleus and depicted as % increase versus control where control was set to 0. Data represent mean and SEM.

**Fig. 6** Effect of Hcy on ATP levels. Increase or decrease in ATP levels (as compared to control levels) in H9c2 cells after incubation with different concentrations of t,t,-Hcy during 24 h. Data represent mean and SEM of four independent experiments.

Indicated by single annexin v positivity. We did not find caspase-3 activation at this concentration. Although annexin V positivity of cells has generally been considered as a marker for apoptosis, in a previous study however, we have demonstrated that this is not absolute as such and that membrane phospholipid flip-flop can be reversible [23], which is supported in this study by the observation that complete inhibition of caspases by Z-VAD-FMK did not affect this phenotype. The caspase-dependent pathway of apoptosis was, therefore, not yet activated at this concentration of Hcy, probably related to the lack of nuclear nitrotyrosin residues.

The highest concentration of 2.7 mM Hcy entailed a completion of the thread-to-grain transition of the mitochondrial reticulum, a further decrease of ATP, translocation of NOX2 to the nucleus and nuclear nitrotyrosin formation, caspase-3 activation, and also a clear increase in
Fig. 7 Effect of Hcy on mitochondrial membrane potential: H9c2 cells were incubated with or without Hcy for 24 h and analyzed with digital imaging microscopy. Green indicates the JC-1 monomers. Red indicates JC-1 aggregates, which are formed in the mitochondria when a sufficiently high membrane potential is reached. (A) H9c2 cells incubated in growth medium alone. Normal heterogeneous population of elongated and round mitochondria have a red fluorescence signal. (B) H9c2 cells incubated with 0.1 mM L-Hcy. An increase in actively respiring mitochondria (red) is visible, and a mostly elongated morphology can be seen. (C) 1.1 mM L-Hcy shows a similar amount of functioning mitochondria in the cells that did not receive Hcy; only the morphology of the mitochondria is mostly round, a phenomenon which is known as thread-to-grain transition. (D) H9c2 cells incubated with 2.7 mM L-Hcy. Some cells have no functioning mitochondria left, but overall there is no significant loss of red signal. The thread-to-grain transition of the mitochondrial reticulum is complete. (E) Analysis of sum intensity in Cy3 signal (red) per condition as an indicator for ΔΨm. Data represent mean and SEM (n = 3).

both single-annexin-V and annexin-V/PI-positive cells. These changes represent a cellular phenotype irreversibly committed to cell death, in which nuclear ROS production apparently plays a deciding role as indicated by the presence of nitrotyrosin residues. Whether cell death then follows the process of apoptosis or necrosis depends on the remaining concentration of ATP. Apoptosis is energy consuming, and therefore depends on minimal residual ATP concentrations, whereas necrosis is the default route in the absence of ATP [39].

Induction of apoptosis generally follows one of two pathways. The first pathway, the death receptor or extrinsic pathway, is activated by ligand binding of cell-surface-expressed death receptors. The second pathway, the mitochondrial or intrinsic pathway, is promoted by physical or chemical stress agents [40]. Intracellular accumulation of Hcy and its metabolites constitutes considerable intracellular stress [15, 17, 41]. In line with this notion, several studies in endothelial cells have suggested that Hcy-induced apoptosis follows the mitochondrial pathway [42,43]. Our study in cardiomyocytes has shown that ΔΨm is modified in a concentration-dependent manner, and thereby supports the concept that Hcy-induced apoptosis mainly follows the mitochondrial pathway, also in cardiomyocytes. However, in addition we found that translocation of NOX2 to the nucleus coinciding with the presence of nitrotyrosin residues indicative for ROS production is even more important in the effector phase of the process of apoptosis inducer in cardiomyocytes.

Since we only found a significant decrease in L-Hcy at the highest concentrations, but not with 0.1 mM α-L-Hcy we also determined SAH concentrations. Several studies
provided further evidence that SAH could be the culprit in Hcy-mediated damage since it is a potent inhibitor of various methylation reactions [32–34]. In this study we found that only the concentration of 2.7 mM \( \text{d,L}-\text{Hcy} \) resulted in a significant increase in SAH, which was the concentration that resulted in cell death of cardiomyocytes. Therefore our study underlines that SAH is an important mediator in the induction of cell death of cardiomyocytes by Hcy.

A limitation of the present study is the use of relatively high concentrations of Hcy. However, it should be noted that cells in culture usually up-regulate their Hcy-metabolizing enzymes, in particular cystathionine beta-synthase (CBS) [44]. Furthermore short-term exposure to high Hcy concentrations may reflect a prolonged exposure to moderately elevated Hcy concentrations as occur life-long in patients. Therefore, our data may provide, at least in part, an explanation why HHC contributes to heart failure.

**Conclusion**

Hcy is increasingly recognized as an independent cardiovascular risk factor [5–8] and an extensive literature exists that links Hcy-induced changes in endothelial [15, 26, 45–48] and smooth muscle cells [16, 17, 49–52] to atherosclerotic disease. The Framingham Heart Study also showed that increased plasma Hcy levels are a risk factor
for the development of heart failure [20]. To our knowledge, no reports have yet been published on the effect of Hcy on cardiomyocytes specifically. Our study now clearly indicates that Hcy, and possible SAH, indeed affects also cardiomyocytes, primarily by inducing apoptosis/necrosis in a concentration-dependent manner, in which nuclear ROS production, related to nuclear NOX expression, is crucial.

Acknowledgement This project is financed by the Dutch Heart Foundation (grant 2002B010) and the iCART-VU (nr. 200370)

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CONCENTRATION-DEPENDENT EFFECTS OF HOMOCYSTEINE ON CARDIOMYOCYTE VIABILITY