

Mechanisms of Hg species induced toxicity in cultured human astrocytes: genotoxicity and DNA-damage response

Cite this: *Metallomics*, 2014, 6, 662

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The toxicologically most relevant mercury (Hg) species for human exposure is methylmercury (MeHg). Thiomersal is a common preservative used in some vaccine formulations. The aim of this study is to get further mechanistic insight into the yet not fully understood neurotoxic modes of action of organic Hg species. Mercury species investigated include MeHgCl and thiomersal. Additionally HgCl₂ was studied, since in the brain mercuric Hg can be formed by dealkylation of the organic species. As a cellular system astrocytes were used. *In vivo* astrocytes provide the environment necessary for neuronal function. In the present study, cytotoxic effects of the respective mercuricals increased with rising alkylation level and correlated with their cellular bioavailability. Further experiments revealed for all species at subcytotoxic concentrations no induction of DNA strand breaks, whereas all species massively increased H₂O₂-induced DNA strand breaks. This co-genotoxic effect is likely due to a disturbance of the cellular DNA damage response. Thus, at nanomolar, sub-cytotoxic concentrations, all three mercury species strongly disturbed poly(ADP-ribosyl)ation, a signalling reaction induced by DNA strand breaks. Interestingly, the molecular mechanism behind this inhibition seems to be different for the species. Since chronic PARP-1 inhibition is also discussed to sacrifice neurogenesis and learning abilities, further experiments on neurons and *in vivo* studies could be helpful to clarify whether the inhibition of poly(ADP-ribosyl)ation contributes to organic Hg induced neurotoxicity.

Received 12th November 2013,
Accepted 3rd February 2014

DOI: 10.1039/c3mt00337j

www.rsc.org/metallomics

Introduction

The three most exposure relevant chemical forms of mercury (Hg) are elemental Hg, mercuric Hg and organometallic compounds. Among these methylmercury is by far the most common in the environment and in the aquatic food-chain. In addition to the consumption of inorganic Hg contaminated food, inorganic Hg exposure might occur through medicinal products.¹ Inhaled elemental Hg vapour from dental amalgam is another source that is likely to increase internal Hg exposure.² Methylmercury exposure occurs nearly exclusively *via* fish and seafood, with generally 80–100% of total fish Hg being

methylmercury.² In Hg polluted areas in China, methylmercury contaminated rice is a further possible contributor.³ Non-dietary exposure to organic Hg might result from the application of thiomersal (sodium 2-ethylmercurothio-salicylate). Thiomersal is used as a preservative in multidose vials of some vaccines,⁴ as well as in several cosmetic products and cleaning solutions for contact lenses.⁵ Its antimicrobial effect is based on its decomposition in aqueous medium to thiosalicylic acid and the ethylmercury cation. Rapid hydrolysis of thiomersal in aqueous biological solution has been demonstrated before.⁶

In 2003, the Joint FAO/WHO Expert Committee on Food Additives (JECFA) revised the PTWI for methylmercury to 1.6 µg kg⁻¹ body weight (b.w.); developmental neurotoxicity was identified as the most sensitive toxicological endpoint. In 2006 JECFA confirmed this PTWI.⁷ Based on the information that beneficial nutrients in fish may have confounded previous adverse outcomes in some of these studies, the European Food Safety Authority (EFSA) Scientific Panel on Contaminants in the Food Chain established in December 2012 a TWI for methylmercury of 1.3 µg kg⁻¹ b.w. Moreover, the Panel concluded that high fish consumers, which might include pregnant women,

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may exceed this TWI by up to six-fold.² Regarding inorganic Hg, in line with JECFA,¹ the EFSA Panel has established a TWI of $4 \mu\text{g kg}^{-1}$ b.w. The estimates of average dietary inorganic Hg exposure in Europe do not exceed the TWI.²

In general mechanisms of toxic effects of mercuric Hg, methylmercury and thiomersal are believed to be similar. Relative toxicities of the respective species are strongly related to their bioaccessibility, accumulation and eventually metabolism or transformation to other Hg species in sensitive tissues. Based on limited lipophilicity mercuric Hg is unlikely to readily cross the blood–brain or the blood–CSF barrier. Nephrotoxicity is the most sensitive endpoint following chronic ingestion of inorganic mercury.^{1,2} In contrast organic Hg species are able to cross these barriers, allowing accumulation in the brain and neurotoxic outcomes.⁸

To date the underlying mechanisms of mercury species induced neurotoxicity are poorly understood. On the cellular level multiple modes of action are discussed to contribute, including the induction of oxidative stress, disturbances in calcium homeostasis, disruption of neurotransmitter systems and cytoskeletal alterations.⁹ With respect to genotoxicity, studies in various cultured mammalian cells have shown a genotoxic potential of the three mercuricals. However genotoxic effects arose often only at quite high, already cytotoxic concentrations.^{2–9a}

This study aims to gain further insights into the molecular mechanisms behind mercury species induced toxicity in human brain cells. Thus, toxic effects of methylmercury (in the form of methylmercury chloride) and thiomersal are compared in cultured human astrocytes. In the brain mercuric mercury can be formed by *in situ* dealkylation of the organic Hg species¹⁰ as well as oxidation of elemental Hg.¹¹ Therefore, cellular toxicity of mercuric Hg (in the form of HgCl_2) is characterized in parallel. Endpoints studied include cytotoxicity, bioavailability, genotoxicity and co-genotoxicity, as well as effects on the cellular DNA damage response.

Materials and methods

Preparation of mercury species stock solutions

HgCl_2 (>99.999% purity, Sigma-Aldrich, Deisenhofen, Germany), MeHgCl (>99.9% purity, Sigma-Aldrich) and thiomersal (>97% purity, Sigma-Aldrich) stock solutions in sterile distilled water were prepared shortly before each experiment.

Cell culture and incubation with the test compounds

Astrocytic cultures (CCF-STTG1 (CCL-185TM)) obtained from the American Type Culture Collection (Bethesda, MD, USA) were used as *in vitro* model systems. CCF-STTG1 cells were cultured in RPMI 1640 (Biochrom, Berlin, Germany), supplemented with 10% FCS (PAA Laboratories, Pasching, Austria), 1.4 mM L-glutamine (Biochrom), 100 U mL^{-1} penicillin and 100 mg mL^{-1} streptomycin (PAA), under human cell culture standard conditions at 37 °C with 5% CO_2 in air and 100% humidity. Logarithmically growing cells were treated with the mercury species as described for the respective experiments.

Cytotoxicity testing

Cytotoxicity of the mercury species was elucidated after 24 and 48 h incubation, by quantifying their effect on cell number and lysosomal integrity. Briefly, after 24 or 48 h of incubation with the respective mercuricals, cells were washed with phosphate buffered saline (PBS), trypsinized and cell number was measured using an automatic cell counter CASY-TTC (Roche Innovatis AG, Germany).¹² These measurements are based on noninvasive, dye-free electrical current exclusion with signal evaluation *via* pulse area analysis. Lysosomal integrity was assessed by the neutral red uptake test,¹³ which represents a well-accepted test system to assess cell viability.

Cellular bioavailability

Mercury levels in CCF-STTG1 cells were measured after 48 h incubation with the respective species by cold vapour atomic fluorescence spectrometry (CV-AFS, mercur Plus, Analytik Jena) or inductively coupled plasma mass spectrometry (ICP-MS, iCAP Qc, Thermo Fisher Scientific), as indicated in the respective figures. Additionally, cellular bioavailability was quantified by ICP-MS after 2 and 24 h incubation with the mercury species. Briefly, the mercury species exposed cells were washed with PBS and trypsinized, collected by centrifugation, washed with ice-cold PBS and the cell number as well as cell volume were measured by an automatic cell counter in each sample. After digestion of cells with 65% HNO_3 /30% H_2O_2 (1/1) at 95 °C as described before¹⁴ total Hg was quantified by CV-AFS or ICP-MS.

Generation of DNA strand breaks

DNA strand breaks were determined by the alkaline unwinding technique.¹⁵ Logarithmically growing CCF-STTG1 cells were exposed to the respective mercury species for 2, 24 or 48 h respectively. In the case of combination experiments with H_2O_2 , after preincubation (48 h) with the mercury species, cells were coincubated with 60 μM H_2O_2 for 5 min. Thereafter the culture medium was removed, cells were washed with cold PBS and lesions were quantified and calculated as described before.¹⁵

Cellular levels of poly(ADP-ribosyl)ation

Cellular poly(ADP-ribosyl)ation was measured as described before.¹⁵ Briefly, CCF-STTG1 cells were seeded on Alcian blue coated glass coverslips, cultured for 48 h and incubated with the respective mercury species for 2, 24 or 48 h. Poly(ADP-ribosyl)ation was stimulated by 10 min incubation with 250 μM H_2O_2 . Subsequently, cells were rinsed with PBS and fixed with ice cold methanol. After fixation, cells were rinsed twice with Tris-buffered saline (TBS) and blocked in TBS/0.3% Tween20/1% BSA (Roth) (TTB) at 30 °C for 30 min. Incubation with the primary anti poly(ADP-ribose) antibody 10H¹⁶ was carried out in blocking solution (1 : 300) at 30 °C for 45 min, followed by repeated washing steps with TBS containing 0.3% Tween20. The secondary, Alexa 488-conjugated anti-mouse antibody (Invitrogen) (dilution 1 : 250 in TTB) was applied accordingly. Coverslips were washed again and embedded in Vectashield mounting medium containing 1 $\mu\text{g mL}^{-1}$ DAPI (Vector Laboratories Inc., Burlingame, CA, USA).



Fluorescence analyses were performed applying a Zeiss Axio ImagerM2 wide field fluorescence microscope (Zeiss, Göttingen, Germany). At least 300 cell nuclei per slide were selected by DAPI staining. In the selected areas the relative Alexa-488 fluorescence intensities were quantified using Axio Vision (Version 4.5) imaging software (Zeiss).

Relative *PARP-1*, *PARG*, *LigIII α* and *XRCC1* gene expression

Real time RT-PCR was performed for quantification of *PARP-1*, *PARG*, *LigIII α* and *XRCC1* mRNA levels. Briefly, 48 h after seeding of 2×10^6 astrocytes, cells were incubated with the respective mercury species for 24 h. All further steps were performed exactly as described before.¹⁵ The respective forward (fwd) and reverse (rev) primers *PARP-1* for 5'-CTTGGCCTG CACACTGTCTG-3', *PARP-1* rev 5'-GCAGCGACTCTCAGATCCTG-3' (efficiency 101.3%); *PARG* for 5'-CCTACTGGGTGGTGACATT-3', *PARG* rev 5'-CGTAAGTGACATGCAATCGT-3' (efficiency 92.1%); *LigIII α* for 5'-GATCACGTGCCACCTACCTT-3'; *LigIII α* rev 5'-GGC ATGGACTGTGGTCATGAG-3' (efficiency 96.9%); *XRCC1* for 5'-CGGATGAGAACACGGACAGT-3'; *XRCC1* rev 5'-CCCCGTAAA GAAAGAAGTGC-3' (efficiency 97.4%); β -Actin for 5'-TGGCATCC ACGAACTACC-3', β -Actin rev 5'-GTGTTGGCGTACAGGTCTT-3' (efficiency 80.8%) were applied. The thermal cycling program consisted of the following steps: 1.5 min at 95 °C to activate polymerase, 40 cycles of 30 s for denaturation at 95 °C, 1 min at 60 °C for annealing and elongation and final denaturation for 15 s at 72 °C. β -Actin expression was not significantly changed by the mercury species in the observed concentration range and thus can be used in this approach as an adequate reference gene.

Activity of recombinant PARP-1

Activity of recombinant PARP-1 was quantified by an established immuno-slot-blot technique.^{15,17} Briefly, after 2 min preincubation of PARP-1 (0.69 ng μL^{-1} (61 nM)) with the respective mercury species in preincubation buffer at room temperature, the PARP-1 reaction was carried out for 5 min at 37 °C in reaction buffer. Poly(ADP-ribosyl)ation was stopped by trichloroacetic acid (TCA). After transferring respective aliquots to a PVDF membrane by a slot blotter, washing and blocking of the membrane, PAR polymers were detected by immunoblot analysis using a monoclonal PAR-antibody (10H) (1:1000, in blocking solution at RT for 1 h) and a HRP-conjugated secondary antibody (1:1000 in blocking solution at RT for 1 h). Immunoreactive bands were detected by chemiluminescence using ECL prime detection reagents (GE Healthcare) and a chemiluminescence imaging system (ChemiDocTM XRS, Bio-Rad, Munich, Germany).

Statistical analysis

All experiments were carried out three times at three different days, with at least three independent measurements. Using the raw data, mean standard deviation (SD) was calculated and a statistical analysis was performed by using ANOVAOneWay and ANOVATwoWay tests. As indicated in the respective figure legends significance levels are * $p < 0.05$, ** $p < 0.01$ and *** $p < 0.001$.

Results

Cytotoxicity in cultured astrocytes

HgCl₂ (Fig. 1A) exerted cytotoxic effects at about 10 times higher concentrations as compared to the organic mercuricals (Fig. 1B and C). Cytotoxicity of HgCl₂ increased with incubation time. After 48 h incubation the endpoint lysosomal integrity was significantly more sensitive than effects on cell number. In the case of MeHgCl and thiomersal, after 48 h incubation cell number was significantly more sensitive than lysosomal integrity. This indicates

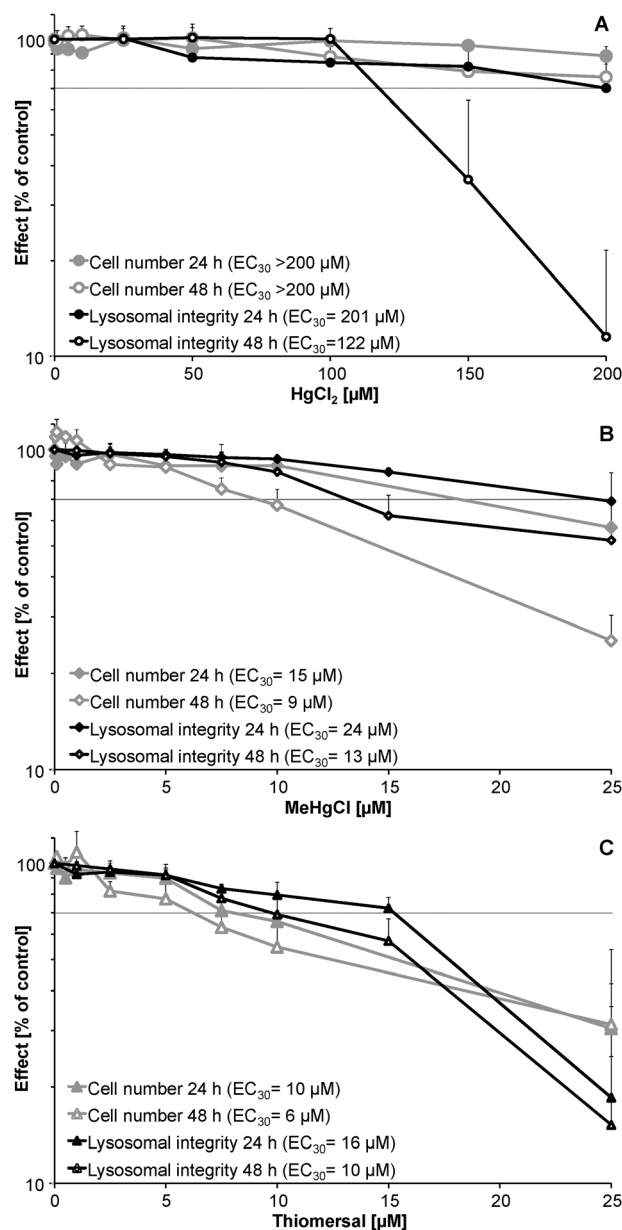


Fig. 1 Cytotoxicity of HgCl₂ [A], MeHgCl [B] and thiomersal [C] in CCF-STTG1 cells after 24 or 48 h incubation. Cytotoxicity was determined by a decrease in cell number and lysosomal integrity. Shown are mean values of at least 3 (cell number) or 18 (lysosomal integrity) determinations + SD. Statistical analysis of the data sets of the different cytotoxicity endpoints (48 h) by two-way ANOVA [A] *** [B] *** and [C] not significant.

that the respective organic mercuricals and HgCl_2 exert their cellular toxicity *via* different toxic modes of action. Time dependency of cytotoxic effects was clearly visible for MeHgCl . In contrast cytotoxic effects of thiomersal did not significantly differ after 24 and 48 h incubation. Thus, after 24 h incubation thiomersal was the most toxic mercurical investigated.

Bioavailability

Cellular bioavailability of the respective mercury species was assessed in a time and concentration dependent manner. After 48 h incubation high cellular mercury levels were quantified *via* CV-AFS. Due to the lower limit of quantification of the applied ICP-MS method (Table 1) low cellular mercury levels were quantified by ICP-MS. As indicated in Fig. 2A–F all applied mercuricals are bioavailable to human astrocytes in a concentration dependent manner. The cellular mercury levels assessed by the two techniques fit well together.

In the case of incubation with HgCl_2 , cellular mercury levels increased in a time dependent manner. This is in strong contrast to the organic mercuricals (Fig. 2A–F).

Determination of DNA strand breaks after mercury species incubation

After 2, 24 and 48 h incubation with subcytotoxic concentrations, HgCl_2 , MeHgCl and thiomersal did not significantly induce DNA strand breaks in CCF-STTG1 cells (Fig. 3A–C). In the case of highly cytotoxic MeHgCl and thiomersal concentrations (25 μM), a massive amount of DNA strand breaks (up to 13 000 lesions per cell) occurred after 48 h of incubation.

Effects of the Hg species on H_2O_2 -induced DNA strand break formation

After 48 h preincubation with the respective mercuricals a concentration dependent increase in DNA strand breaks after combined treatment with 60 μM H_2O_2 was observed, when compared to 60 μM H_2O_2 alone (Fig. 4A–C).

Effects on cellular poly(ADP-ribosyl)ation

After short-term (2 h) and long-term (24, 48 h) incubation the mercury species exerted no effects on poly(ADP-ribosyl)ation in non-stimulated CCF-STTG1 cells (data not shown). Since in

unstressed cells the presence of poly(ADP-ribose) is in general quite low, in the next set of experiments cellular poly(ADP-ribosyl)ation was stimulated by H_2O_2 , to study the impact of the Hg species on this DNA damage related signaling reaction.

In CCF-STTG1 cells a 2–48 h preincubation with subcytotoxic concentrations of HgCl_2 (≥ 0.5 μM), MeHgCl (≥ 0.05 μM) and thiomersal (≥ 0.05 μM) significantly decreased the extent of H_2O_2 -induced poly(ADP-ribosyl)ation (Fig. 5A–C). Inhibition was independent of incubation time with the respective mercuricals.

Effects on *PARP-1*, *PARG*, *LigIII α* and *XRCC1* gene expression

24 h incubation with subcytotoxic concentrations of HgCl_2 (≥ 0.5 μM) led to significantly down-regulated *LigIII α* gene expression; but did not significantly alter *XRCC1* and *PARG* mRNA levels (Table 2). *PARP-1* gene expression was significantly lowered only by the highest concentration of HgCl_2 applied (50 μM). Likewise HgCl_2 , the organic mercuricals did not significantly disturb *PARG* expression (Table 2). *PARP-1*, *XRCC1* and *LigIII α* expression were decreased by cytotoxic concentrations of MeHgCl (10 μM) and thiomersal (10 μM). Additionally, MeHgCl significantly diminished *LigIII α* expression at 1 μM .

Effects on poly(ADP-ribosyl)ation of recombinant *PARP-1*

2 min preincubation of recombinant *PARP-1* with HgCl_2 (≥ 2 μM), MeHgCl (≥ 18 μM) or thiomersal (≥ 200 μM) did significantly diminish *PARP-1* activity (Fig. 6A–C).

Discussion

To further understand the mechanisms behind organic mercury induced neurotoxicity, this study investigated and compared cytotoxic as well as direct and indirect genotoxic effects of HgCl_2 , MeHgCl and thiomersal in human astrocytes.

Astrocytes and microglia represent the two major cell types, which protect optimal brain function; astrocytes occupy 25% to 50% of the brain volume.¹⁸ Furthermore, evidence suggests the essential nature of interactions between neurons and astrocytes for brain function.¹⁹ Therefore, the present study was carried out in human astrocytes.

Organic mercury species are well known to cross the brain regulating barriers and subsequently accumulate in different brain areas, including the cerebellum, brain cortex and retina.²⁰ Even though mercuric mercury cannot pass these barriers, it is present in the brain. Therefore mercuric mercury has to be taken into account, when searching for the mechanisms of mercury species induced neurotoxicity. Thus, inorganic mercury can be formed inside the brain by partial dealkylation of the organic species as well as by oxidation of $\text{Hg}^{0,2,10,11}$.

In the astrocytes the cytotoxicity of the mercuricals increased with rising alkylation level and fit nicely to the cellular bioavailability of the mercuricals. A similar cytotoxic order has been shown recently for these three mercury species in human T leukemic Jurkat cells.⁸ As possible underlying mechanisms for cytotoxicity, several end-points are discussed, including a disturbance of proliferation,²¹ a disruption of cellular stress, redox homeostasis²² and apoptosis.²³

Table 1 Conditions for CV-AFS and ICP-MS analysis of total mercury

Parameter	AFS	ICP-MS
Power	725 W	1550 W
Nebulizer gas		1.075 L min ⁻¹
Cell gas		8% H_2 in He
Cell gas flow		5.11 L min ⁻¹
Isotope		²⁰⁰ Hg
Internal standard		50 ng L ⁻¹ Bi
LOD	1.2 ng L ⁻¹	0.2 ng L ⁻¹
LOQ	3.5 ng L ⁻¹	0.8 ng L ⁻¹
Reducing agent	2% (w/v) SnCl_2 ; 10% (v/v) HCl	
Jetting agent	1% (v/v) HNO_3 ; 2% (v/v) HCl	
Reductant flow	4 mL min ⁻¹	
Sample speed	10 mL min ⁻¹	
Argon flow	170 mL min ⁻¹	



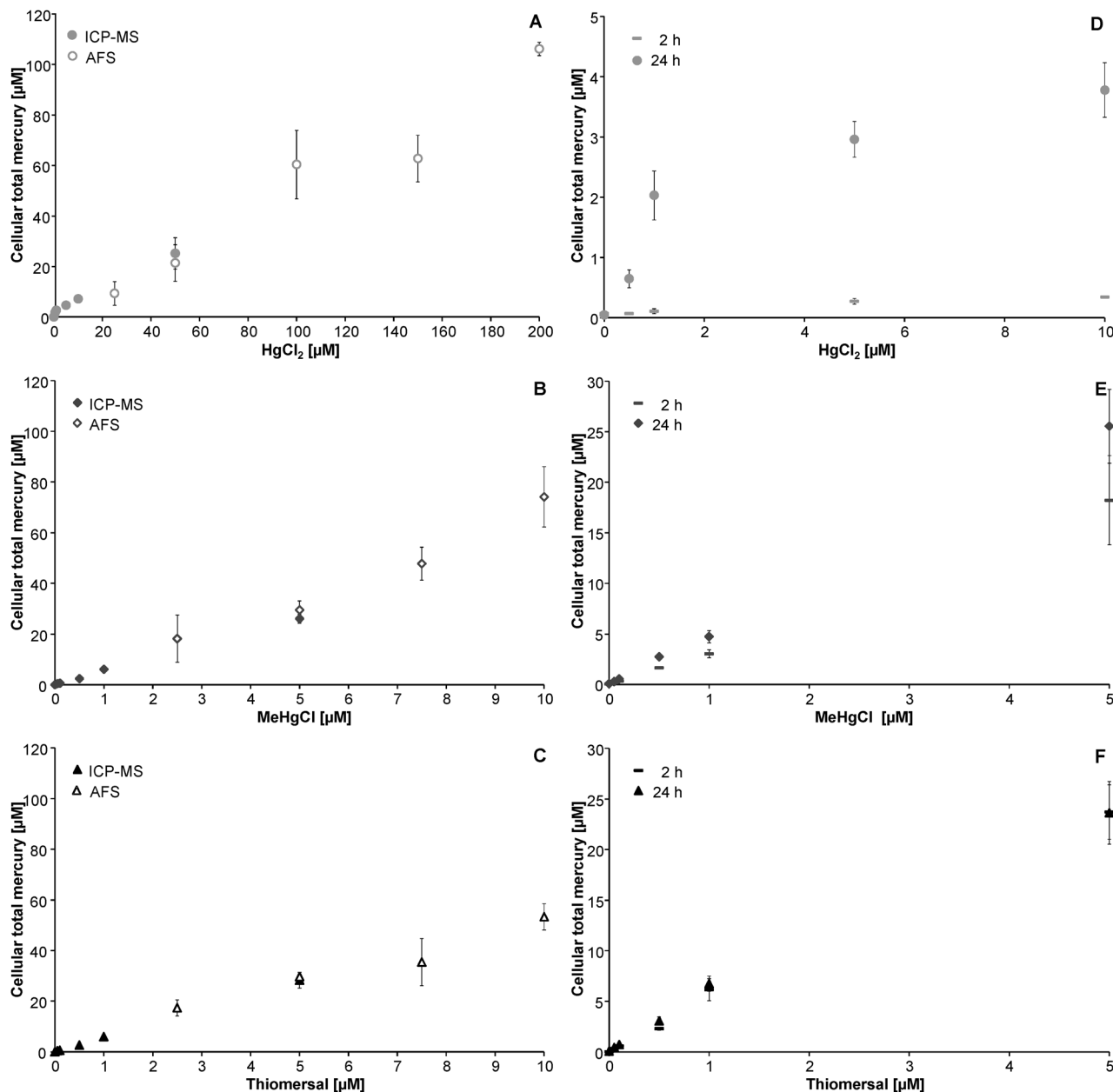


Fig. 2 Cellular bioavailability of the mercury species. Concentration dependent cellular bioavailability after 48 h incubation with HgCl₂ [A], MeHgCl [B] or thiomerseal [C] as determined by AFS (open symbols) and ICP-MS (closed symbols). Time dependent cellular bioavailability of the mercury species within 2–24 h incubation as quantified by ICP-MS [D–F]. Shown are mean values of at least three independent determinations ± SD.

In the present study, lysosomal integrity was affected after HgCl₂ incubation at lower concentrations than cell number. Thus, in the applied astrocytes mercuric Hg seems to disturb the cellular energy level resulting in a collapse of the proton gradient of the lysosomes. In contrast, the two organic mercuricals affected especially the endpoint cell number, which indicates a disturbance of proliferation. Accordingly, cytogenetic studies demonstrated that both organic mercuricals strongly increased the number of binucleated cells in the respective concentration range (data not shown).

In the cytotoxic concentration range cytotoxicity of the mercuricals was accompanied by a massive increase of DNA strand breaks. Nevertheless, in the sub-cytotoxic concentration range

neither HgCl₂, MeHgCl nor thiomerseal induced DNA strand breaks. Thus, it seems unlikely that the mercuricals induce cytotoxicity *via* DNA strand break induction. It is more likely that the massive increase of DNA strand breaks is a consequence of the cytotoxic process, which has been postulated before.²⁴

Whereas the mercury species failed to increase DNA strand breaks in the sub-cytotoxic concentration range, they strongly increased H₂O₂-induced strand breaks. Our data clearly indicate that this co-genotoxic effect is due to a disturbance of the cellular DNA damage response.

One of the first nuclear reactions following DNA strand breaks induction is poly(ADP-ribosyl)ation of proteins. The respective



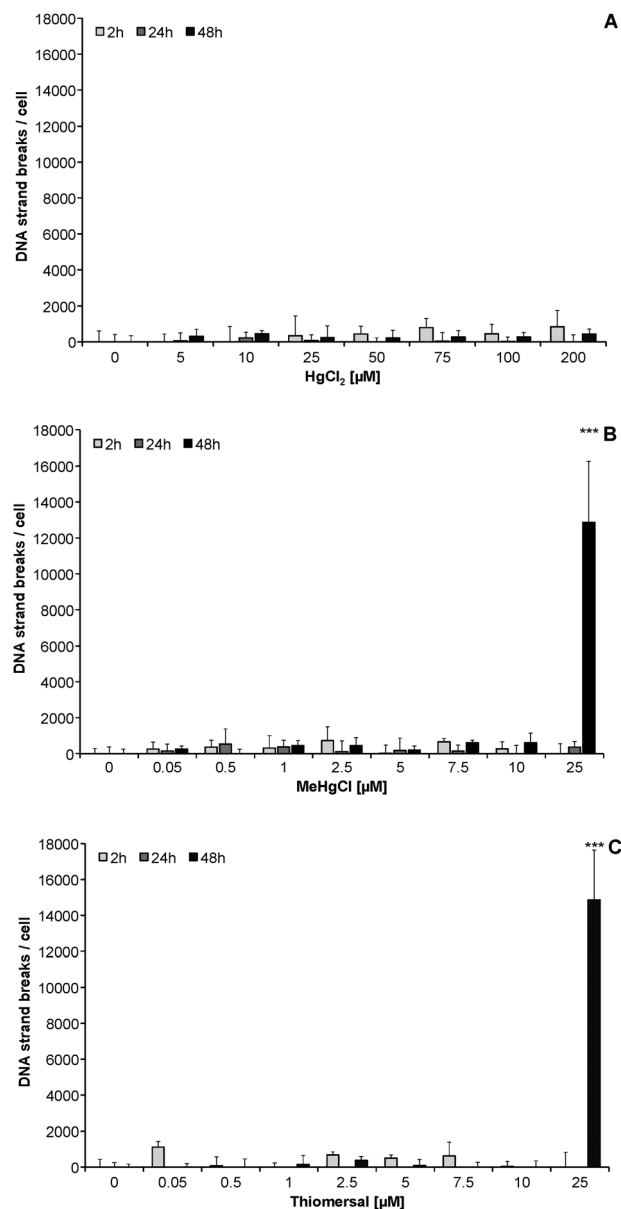


Fig. 3 Number of DNA strand breaks after 2, 24 or 48 h incubation with HgCl_2 [A], MeHgCl [B] or thiomersal [C]. Shown are mean values of at least two independent experiments with three determinations each + SD.

polymerase PARP-1 is responsible for about 90% of cellular poly(ADP-ribosylation).²⁵ The negative charges of poly(ADP-ribose) are believed to cause relaxation of chromatin structure and thereby increase accessibility for other repair proteins, to carry out the repair of the DNA single strand breaks.²⁶ Poly(ADP-ribosylation) of PARP-1 additionally triggers the recruitment of XRCC1, which coordinates and stimulates repair of DNA strand breaks. In the further process $\text{LIGIII}\alpha$, in association with XRCC1, efficiently ligates the respective DNA nicks.²⁷ Thereby, $\text{LIGIII}\alpha$ activity requires the presence of poly(ADP-ribosylated) PARP-1.²⁸

In this study poly(ADP-ribosylation) is identified for the first time as a novel, highly sensitive endpoint in the toxic mode of action of mercury species. For all three mercuricals a strong inhibition of this essential DNA damage response related

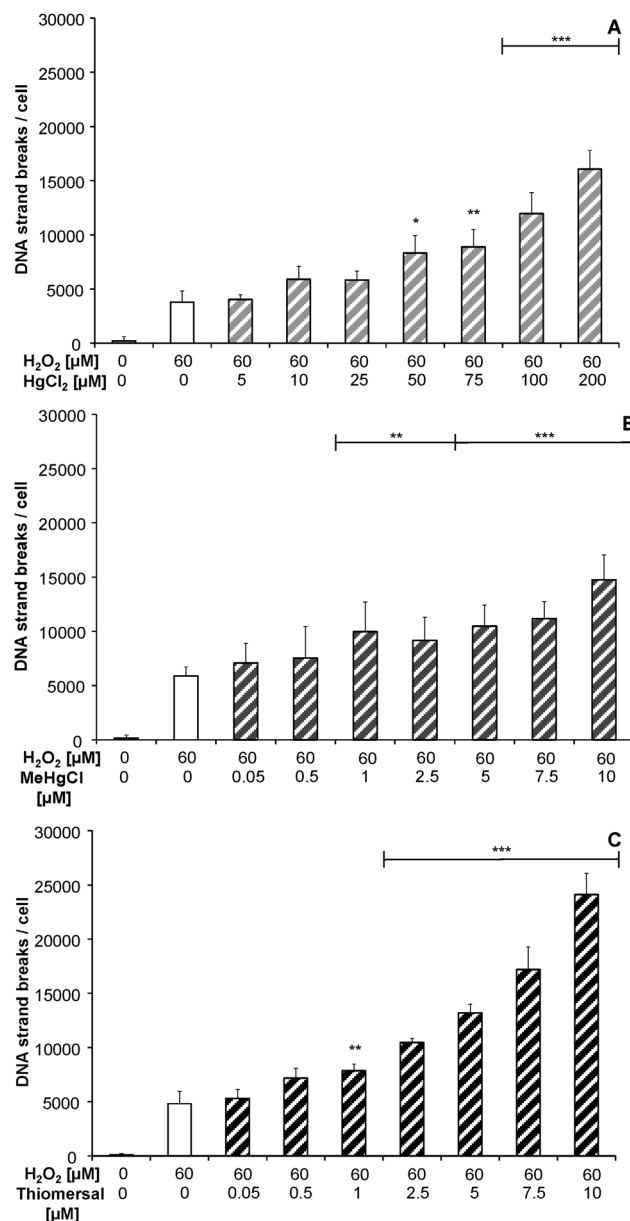


Fig. 4 Impact of a 48 h preincubation with HgCl_2 [A], MeHgCl [B] or thiomersal [C] on the number of DNA strand breaks induced by hydrogen peroxide (5 min, 60 μM) in CCF-STTG1 cells. Displayed are mean values of at least two independent experiments with three determinations each + SD.

signaling reaction was detected after 2–48 h incubation with subcytotoxic concentrations of the mercury species. A significant inhibition of H_2O_2 -induced poly(ADP-ribosylation) was observed by 0.5 μM HgCl_2 and 0.05 μM MeHgCl or thiomersal. These incubation concentrations resulted after 2 h incubation in cellular mercury levels of 0.04, 0.20 and 0.36 μM , respectively.

As underlying mechanism for the inhibition, we could exclude a mercury species induced delay of the signalling reaction process, by carrying out time-dependent poly(ADP-ribosylation) experiments (data not shown). The inhibition was not due to a decrease of *PARP-1* gene expression. A decrease of the *PARP-1* substrate NAD^+ could also be excluded, especially since the observed inhibition

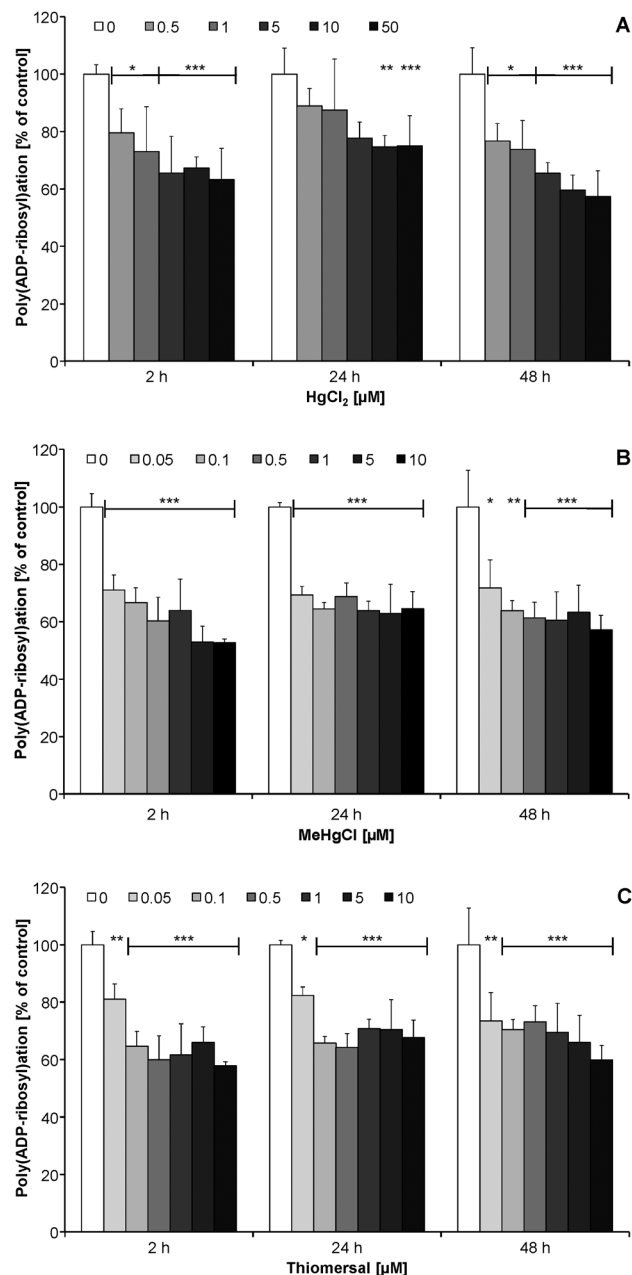


Fig. 5 Induction of poly(ADP-ribose)ylation by H_2O_2 and inhibitory effect of 2, 24 and 48 h preincubation with HgCl_2 [A], MeHgCl [B] or thiomersal [C]. Logarithmically growing CCF-STTG1 cells were preincubated with the mercuricals for 2, 24 and 48 h and treated with $250 \mu\text{M}$ H_2O_2 for 10 min in the continued presence of the mercury species. Shown are mean values of at least three independent determinations with at least 200 cells each + SD.

of poly(ADP-ribose)ylation occurred in absolutely non-cytotoxic concentrations. None of the mercury species increased gene expression of poly(ADP-ribose)glycohydrolase (*PARG*), with the respective protein being responsible for degradation of cellular poly(ADP-ribose).²⁹ We cannot exclude that the mercury species stimulate *PARG* activity. This would result in an increased degradation of cellular poly(ADP-ribose) and thus also in a decrease of cellular poly(ADP-ribose)ylation.

Table 2 Effect of the mercury species on *PARP-1*, *PARG*, *XRCC1* and *LigIII α* gene expression. CCF-STTG1 cells were incubated with HgCl_2 , MeHgCl or thiomersal for 24 h. Relative gene expression was determined by real time RT-PCR; mean values of at least 2 independent determinations with 3 measurements each, referring to the control and normalized to β -Actin $\pm$ SD

C [μM]	<i>PARP-1</i>	<i>PARG</i>	<i>XRCC1</i>	<i>LigIIIα</i>
HgCl_2				
0	1.00 $\pm$ 0.08	1.00 $\pm$ 0.08	1.00 $\pm$ 0.07	1.00 $\pm$ 0.01
0.5	0.97 $\pm$ 0.08	0.95 $\pm$ 0.08	0.87 $\pm$ 0.15	0.89 $\pm$ 0.09*
1	0.91 $\pm$ 0.13	0.85 $\pm$ 0.13	1.04 $\pm$ 0.10	0.72 $\pm$ 0.02***
5	0.92 $\pm$ 0.10	0.84 $\pm$ 0.10	1.01 $\pm$ 0.07	0.69 $\pm$ 0.05***
10	0.90 $\pm$ 0.07	0.84 $\pm$ 0.17	0.95 $\pm$ 0.04	0.68 $\pm$ 0.06***
50	0.75 $\pm$ 0.14***	0.77 $\pm$ 0.21	1.07 $\pm$ 0.07	0.71 $\pm$ 0.06***
MeHgCl				
0	1.00 $\pm$ 0.12	1.00 $\pm$ 0.09	1.00 $\pm$ 0.04	1.00 $\pm$ 0.01
0.05	0.95 $\pm$ 0.08	1.07 $\pm$ 0.07	1.05 $\pm$ 0.06	0.96 $\pm$ 0.15
0.1	0.94 $\pm$ 0.16	1.01 $\pm$ 0.15	1.09 $\pm$ 0.11	0.87 $\pm$ 0.11
0.5	1.03 $\pm$ 0.09	1.09 $\pm$ 0.19	1.06 $\pm$ 0.07	1.01 $\pm$ 0.13
1	1.03 $\pm$ 0.14	1.11 $\pm$ 0.12	1.09 $\pm$ 0.09	0.72 $\pm$ 0.08**
10	0.71 $\pm$ 0.15**	1.20 $\pm$ 0.15	0.78 $\pm$ 0.06***	0.65 $\pm$ 0.18***
Thiomersal				
0	1.00 $\pm$ 0.10	1.00 $\pm$ 0.05	1.03 $\pm$ 0.08	1.00 $\pm$ 0.04
0.05	1.06 $\pm$ 0.09	1.02 $\pm$ 0.08	1.15 $\pm$ 0.13	1.01 $\pm$ 0.09
0.1	1.00 $\pm$ 0.17	0.94 $\pm$ 0.13	1.13 $\pm$ 0.16	0.88 $\pm$ 0.14
0.5	0.95 $\pm$ 0.06	1.00 $\pm$ 0.11	1.00 $\pm$ 0.07	0.85 $\pm$ 0.10
1	0.87 $\pm$ 0.12	0.94 $\pm$ 0.15	1.03 $\pm$ 0.16	0.91 $\pm$ 0.04
10	0.76 $\pm$ 0.13*	1.08 $\pm$ 0.17	0.71 $\pm$ 0.09***	0.69 $\pm$ 0.08***

As possible mechanism for the disturbance of this essential cellular signalling reaction an inhibition of *PARP-1* activity might be possible. All three mercuricals were able to diminish the activity of recombinant *PARP-1*. Here the inhibitory effect decreased with increasing alkylation level of the mercuricals. In the case of HgCl_2 and MeHgCl the low micromolar concentrations, which effectively disturbed recombinant *PARP-1* activity, are likely to be reached in the cellular system. In the astrocytes disturbance might at least partly be due to an interaction of the mercuricals with cellular *PARP-1* activity. One possible mechanism could be a direct interaction of HgCl_2 and MeHgCl with the zinc finger structures³⁰ of *PARP-1*. In the case of thiomersal, the recombinant *PARP-1* protein data reveal that this mechanism is unlikely to be relevant for the cellular system. Nevertheless, thiomersal might inactivate cellular *PARP-1* activity also indirectly, *e.g.* by a generation of reactive oxygen species or also by a metabolism of the mercurical to another more effective mercury species. This might be clarified by cellular mercury speciation studies in the future.

Since MeHgCl and especially HgCl_2 ($\geq 0.5 \mu\text{M}$) downregulate *LigIII α* gene expression, repair of strand breaks might additionally be blocked by a disturbance of the ligation process; thiomersal affected *LigIII α* expression only in a cytotoxic concentration.

In summary, the single mercury species affect common but also different targets of the cellular stress response. Thus, in the case of a mixed mercury species exposure in brain target cells, the DNA damage response will be efficiently disturbed. Regarding the inhibition of the DNA strand break repair, all three mercuricals can disturb the signalling reaction poly(ADP-ribose)ylation.



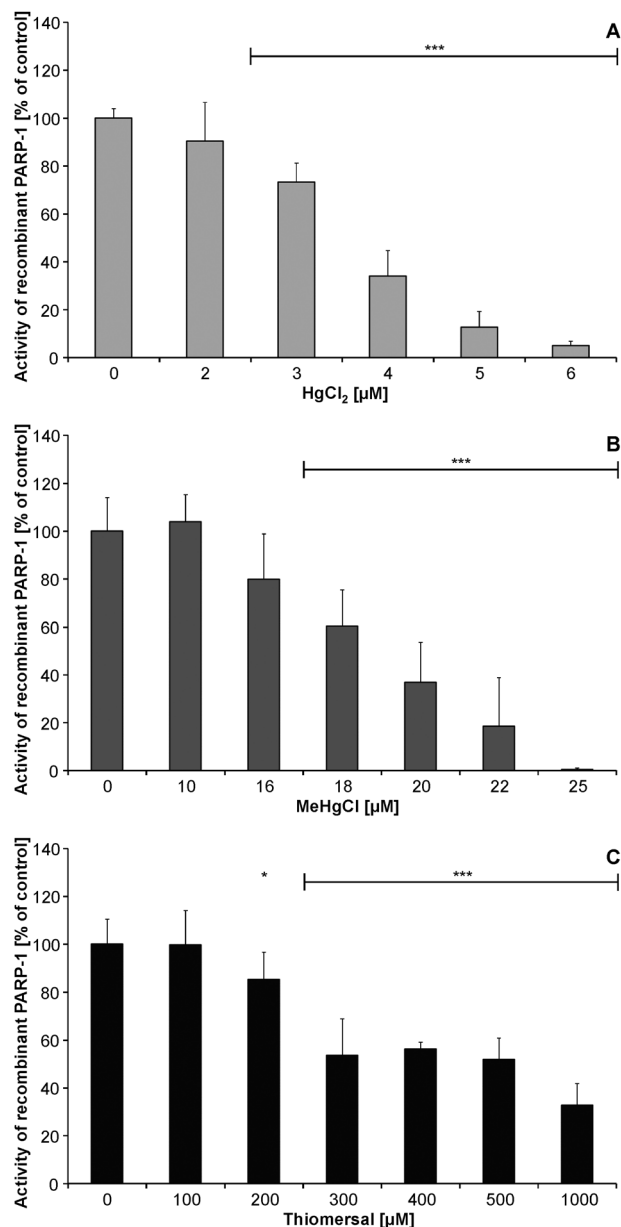


Fig. 6 Impact of HgCl₂ [A], MeHgCl [B] or thimerosal [C] on the activity of recombinant PARP-1. Shown are mean values of at least three independent determinations with three measurements each + SD.

HgCl₂ and MeHgCl downregulate LigIII α , and HgCl₂ can additionally disturb the energy level and thus limit cellular ATP, which is necessary for the ligation process. Poly(ADP-ribosylation) is affected by the respective mercuricals at concentrations as low as 50 nM and most likely by different mechanisms. This endpoint should be further studied in experiments applying mixed mercury species exposure, to rule out if the disturbance of this essential signalling reaction contributes to organic mercury induced neurotoxicity. Hence it is well known that the cellular reduction of DNA damage induced poly(ADP-ribosylation) has serious consequences. DNA strand breaks, but also other DNA lesions, including oxidative base modifications, are no longer efficiently repaired and might accumulate.

Subsequently, astrocytes become dysfunctional and/or undergo cell death. Death of astrocytes can disturb neural development, but also cause death of neurons.³¹ Furthermore, it is possible that the mercuricals disturb damage induced poly(ADP-ribosylation) also in neurons. This has to be confirmed in future studies, especially since abnormal neuronal development or even a loss of neurons seems to play a role in many chronic neurodegenerative diseases such as ischaemia, Alzheimer's, Parkinson's and Huntington's diseases.³² Furthermore, continued PARP-1 inhibition is discussed to result in a disturbance of neurogenesis and learning abilities in perinatal CNS.³³ Whether there is a correlation to the decrement in intelligence and impairment of memory in children highly exposed to mercury species,³⁴ should be tested in further studies.

Acknowledgements

The authors would like to thank Prof. Dr Hans-Joachim Galla (Institute of Biochemistry, University of Münster, Germany) for the CCF-STTG1 cells and PD Dr. Sascha Beneke (Institute of Veterinary Pharmacology and Toxicology, University of Zürich Switzerland) for helping us with the recombinant PARP-1 assay.

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