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New insights on mechanisms underlying methylmercury-induced and manganese-induced neurotoxicity

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Abstract

Toxic and essential elements are widely distributed in the Earth's crust and individuals may be exposed to several of them. Indeed, exposure to toxic elements such as mercury (Hg) can be a potential health risk factor of health, mainly by ingestion of fish containing methylmercury (MeHg). On the other hand, essential elements such as manganese (Mn) play an important role in physiological process in human body. However, Mn overexposure may cause toxic effects. In this respect, the neurotoxic effects of MeHg and Mn on the developing brain are well recognized. Therefore, in this critical review, we address the effects of MeHg and Mn on cell signaling pathways which may contribute to molecular mechanisms involved in MeHg- and Mn-induced neurotoxicity.

Keywords

Mercury; Methylmercury; Manganese; Heavy metal; Signaling pathways; Neurotoxicology

1. Introduction

Exposure to heavy metals and metalloids such as arsenic, lead, cadmium and mercury (Hg) can lead to the development of diseases (1, 2). Yet, some metals are considered “essential elements”. These elements, such as manganese (Mn), selenium and copper, play important roles in the regulation of critical enzyme systems and are essential for several physiological process (3, 4). However, when essential elements are present at higher concentrations, they may disrupt normal biological functions and induce cellular stress responses, thus contributing to development of diseases (5, 6).

Regarding toxic elements, Hg is ranked 3rd in the list of Hazardous Substance Priority List established by the US ATSDR in 2019. This hazardous pollutant occurs in different chemical

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Declaration of interests

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forms: elemental Hg (Hg^0), inorganic Hg compounds (Hg^{2+}), and organic Hg compounds, as methylmercury (MeHg). In aquatic environment, the Hg^{2+} can be biomethylated by aquatic sulfate-reducing bacteria, generating MeHg, which has a substantial biomagnification potential and accumulates along the food chain. Consequently, fish intake is the major source of MeHg exposure in humans (7, 8). Several studies have reported that MeHg exposure may lead to neurological alterations, including cognitive and motor dysfunction, and decreases in memory and learning (9, 10).

The essential element Mn is required to several biological process, including brain and skeletal development, immune response and others (11, 12). Moreover, it acts as an important cofactor for numerous enzymes, participating in synthesis and metabolism of amino acids, proteins, lipids as well as enzymes to defense of the organisms against oxidative stress. The major source of exposure to Mn is through the diet (11). In addition, inhalation of high levels of airborne Mn aerosols in occupational activities represents an important source of exposure to workers. It is well documented that Mn exposure may lead to its excessive accumulation in the central nervous system (CNS), resulting in manganism (13). Furthermore, chronic alterations of Mn levels in the brain may trigger the development of neurodegenerative diseases, such as Alzheimer's disease (AD) and Parkinson's disease (PD) (14, 15).

Currently, the cumulative evidence suggests the involvement of Mn and MeHg in adverse neurological toxic effects. Interestingly, co-exposure to MeHg and Mn causes more pronounced toxic effects than single metal exposure. It was reported that co-exposure to these metals in *Caenorhabditis elegans* (*C. elegans*) caused developmental delays in worms, increased enzymes associated with the antioxidant system and cholinergic degeneration (16). Therefore, both metals could be involved in development of neurotoxicity. However, the molecular mechanisms responsible for metal-induced neurotoxicity cannot be explained by a single process. Rather, several changes in cell signaling pathways involved in the regulation of the cells in the brain contribute to those effects. In this narrative review, we summarize the current knowledge regarding the molecular mechanisms involved in MeHg and Mn neurotoxicity.

2. Methylmercury: new insights in neurotoxicity

Methylmercury (MeHg) is a naturally occurring potent neurotoxin that is produced by microorganisms in water sediment through transformation of inorganic mercury. Human exposure to MeHg is predominantly from consumption of fish that bioaccumulate MeHg from lower tropical organisms in the water. The relatively high level of mercury in top tropical fish poses a potential health risk for people whose regular protein source are marine food, especially developing children and childbearing women. Acute MeHg exposure at high levels causes permanent neuronal damage, while chronic MeHg exposure at developmental stage has long-term impacts on neurobehavioral functions. MeHg-induced neurotoxicity involves multiple mechanisms including oxidative stress, repression of protein translation, disruption of calcium homeostasis and mitochondrial energetics, and post-translational modification of proteins (17). Accumulating evidence shows that these mechanisms are

involved in several important cellular processes and functions, culminating in neuronal toxicity (18–20).

Low-level MeHg exposure at critical developmental stage has profound impact on neurobehavior functions and neuronal cell integrity. Gestational exposure to MeHg has been shown to induce cerebellar synaptic and neuritic remodeling during the perinatal period through modulation of the TrkA pathway and Arc expression (18). Emerging evidence utilizing highly tractable animal models illustrates that cellular morphology and organismal behavioral patterns were altered by low levels of MeHg (21, 22).

By comparing neuronal morphology of animals with various long-term MeHg exposure regimes, a chronic MeHg exposure model with *C. elegans* was recently established for evaluation of dopaminergic (DAergic) neurodegeneration (23). A novel type of neuronal homeostasis was demonstrated in *C. elegans* showing that proteotoxic stress promoted neuronal cell removal of spatially organized proteins and organelles destined for degradation (24). Studies with the *C. elegans* model demonstrated that the removal process was inhibited by MeHg (21, 25). The stress inducible protein 1 (STI-1) is a co-chaperone that assists protein folding in the intermediate stage of the chaperone-assisted protein folding pathway (Hsp70/Hsp40/Hsp90) by transferring client proteins from the early complex to the intermediate complex. Further analysis showed that STI-1 plays a critical modulatory role in MeHg-induced disruption of the regulated removal process (21).

MeHg has long been recognized as a disruptor of protein homeostasis and oxidative balance. Altered functions of membrane transporters and receptors following MeHg exposure are one of underlying mechanisms for neurobehavioral damages. For example, rats with developmental MeHg exposure were sensitive to the effects of the dopamine agonist, d-amphetamine, and the disruption of dopamine neurotransmission by MeHg contributes to the baseline-dependent inhibition of behavior by d-amphetamine (26). Novel mechanistic insights were also derived from studies showing that the cytotoxic effects of MeHg in motor neurons were mediated by the activation of AMPA receptors (27). Furthermore, shifted glutamatergic excitotoxicity by MeHg may contribute neurodegeneration and loss of motor function, given that MeHg induces hyperexcitability in lumbar spinal motor neurons by increasing activity of Ca²⁺-permeable AMPA receptors (28).

In addition to the classical roles of antioxidant pathways in MeHg-induced cytotoxicity, recent studies suggest that the antioxidant signaling network involves several important cellular domains including differentiation and protein degradation (29, 30). In a *Drosophila* model, muscle development defect following MeHg exposure could be rescued by neuron-specific upregulation of one of the central transcriptional factors invoked for antioxidant defense against MeHg toxicity, cap-n-collar C (CncC), the homolog of nuclear factor erythroid 2-related factor 2 (Nrf2) (29). Surprisingly, MeHg-induced lethality could be potentiated by muscle-specific CncC upregulation, while neuron-specific upregulation of CncC was protective (29).

Neuronal NADPH diaphorase (NADPH-d) acts as a cosubstrate for neuronal nitric oxide synthase, which produces the gaseous neuromodulator nitric oxide (NO). The astrocytic

activity of NADPH-d and cell numbers was reduced in the visual cortex of rats with chronic MeHg exposure. However, it seemed that the morphology of NADPH-d neurons was spared from the toxic effects of MeHg. A plausible hypothesis was proposed that the negative effect of chronic MeHg poisoning on both the synthesis and transport of NADPH-d in afferent pathways to the visual cortex contributes to the decrease in astrocytic NADPH-d reactivity (19). In addition, MeHg exposure induced degradation of astrocytic hypoxia-inducible factor-1 α (HIF-1 α) via generation of reactive oxygen species (ROS). Overexpression of HIF-1 α attenuated MeHg-induced cytotoxicity (30). Reactive sulfur species (RSS) contain mobilized sulfur that readily captures xenobiotic electrophiles, forming sulfur adducts. The enzyme necessary for RSS synthesis, cystathionine γ -lyase (CSE), plays a protective role in MeHg-induced motor impairment. CSE-deficient mice were susceptible to toxic effects of MeHg, which can be rescued by restoration of RSS with supplementation of sodium tetrasulfide (31).

It has been recently also recognized that the toxicity of MeHg is modulated by bacteria (32). It was shown that the intestinal microbiota played a significant role in MeHg-induced impairment of locomotor activity. (33). Additionally, metabolomics profiling showed that MeHg induce changes of intestinal microbial composition as well as BDNF level, suggesting a potential link between gut microbiota and MeHg-induced neurotoxicity (20).

3. Manganese: new insights in neurotoxicity

The acute or chronic exposure to Mn may affect important cell signaling pathways that regulate cell survival, differentiation, and apoptosis. Indeed, several factors interplay to form the cascade of events involved in Mn neurotoxicity, such as oxidative stress, neuroinflammation, transporter dysregulation, mitochondrial dysfunction and protein misfolding (34, 35). Moreover, Mn-induced neurotoxicity shares pathways associated with the development of neurodegenerative diseases (14).

Overexposure to Mn may be associated with changes in protein aggregation such as A β and Tau that are AD hallmark. Recently, Wang et al. reported that rats showed increased A β ₁₋₄₀ and Tau production in brain after Mn exposure. The authors also reported that the levels of NLRP3 (nucleotide binding and oligomerization domain-like receptor family pyrin domain-containing 3) inflammasome and inflammatory factors such as IL-1 β , IL-18 were higher in the brain of Mn exposed rats compared to the control, suggesting that Mn promoted the activation of NLRP3 inflammasome, and that the expression of inflammatory factors was upregulated in cerebral tissues (36). Peng et al. (2020) demonstrated that Mn can activate inflammatory pathways such as NF- κ B, causing inflammation by increasing the phosphorylation of p65 and I κ B- α as shown by increased expression of p-p65 and p-I κ B- α in Mn-treated BV2 cells and in the basal ganglia of Mn-exposed rats. In addition, the expression of NLRP3 and cleaved caspase 1 (Cleaved CASP1) was significantly increased after Mn treatment in BV2 cells and in the basal ganglia of rats (37). The effects of Mn exposure on NLRP3 inflammasome and neuroinflammation appear to be associated with mitochondrial dysfunction. Indeed, a recent study has shown that Mn exposure decreased the abundance of mitochondrial fusion protein 2 and Mfn2 degradation. Mn-induced mitochondrial dysfunction stimulated cell-to-cell transfer of the inflammasome adaptor

protein ASC (apoptosis associated speck-like protein containing a CARD) through exosomes, further spreading inflammasome activation (38).

The NF- κ B signaling pathway has an important role in inflammatory processes. Under physiological conditions, NF- κ B is present in the cytosol and bound to I κ B- α , its inhibitory protein. The activation of I κ B- α leads to phosphorylation and degradation of I κ B α , leading to phosphorylation of NF- κ B and its translocation from the cytosol to the nucleus where NF- κ B activates the transcription of target genes such as TNF- α , IL-1 β , and IL-6 (39). In fact, previous studies showed that Mn activated NF- κ B signaling, changing the inflammatory gene expression (40). In agreement, it was reported that Mn stimulated NF- κ B signaling pathways in the hippocampus and striatum of rats, increasing pro-inflammatory cytokines TNF- α , IL-1 β , and IL-6 levels (41). An *in vitro* study showed that Mn activated the NF- κ B pathway in BV2 microglia, further propagating inflammatory cytokine production, concomitant with increased phosphorylation of P65 protein and mRNA expression in BV2 microglia (42). Collectively, these studies support that NF- κ B signaling activation is involved in the Mn-induced neuroinflammatory process.

Phosphatidylinositol 3 kinase (PI3K) is a phosphatidylinositol kinase that contributes to oxidative stress and regulation of cell differentiation, growth and apoptosis (43). The downstream mediator of PI3K is serine-threonine protein kinase (Akt) which acts as antiapoptotic activity by preventing the release of cytochrome C from mitochondria and inactivating fork head box transcription factors (FOX). In addition, Akt can activate or inhibit its downstream target proteins (NF- κ B, caspase-9, Bad, FoxO3a) (43). It has been reported that Mn exposure activated PI3K/Akt signaling pathway in rat hippocampus, leading to inhibition of the transcription function of apoptotic genes such as Bcl-2 and caspase-3, and the activation of Bax, suggesting that apoptosis may underlie the cognitive dysfunction in these animals (44). Moreover, Cheng et al. (2018) demonstrated that mRNA levels of Akt-1 and FoxO3a were decreased after Mn exposure. However, Akt protein levels were increased, indicating that chronic Mn exposure activated the PI3K/Akt pathway via phosphorylation of Akt, in turn, inhibiting the transcription function of apoptotic genes and leading to cell survival (45). Using *C. elegans* model, Peres et al. (2018) showed that worms with loss of Akt (akt-1 and akt-2) was associated with higher resistance to Mn compared to wild-type worms, suggesting that Akt may serve as a potential therapeutic target for Mn neurotoxicity (46).

4. Conclusions

Growing evidence suggests that MeHg and Mn overexposure leads to toxic effects, particularly in the brain, contributing to neurodegenerative diseases. Exposure to heavy metals such as MeHg and Mn is an important public health concern. Overall, the findings summarized in this review (Figure 1) suggest that the understanding of how toxic and sub-toxic levels of Me Hg and Mn may stimulate several critical cells signaling pathways that are involved in variety of biological processes and illness states may be used as a strategy to treat or prevent neurotoxic effects induced by these metals. In addition, research using “omic” tools such as proteomic, transcriptomic and bioinformatic analyses are necessary and afford new directions in toxicological research.

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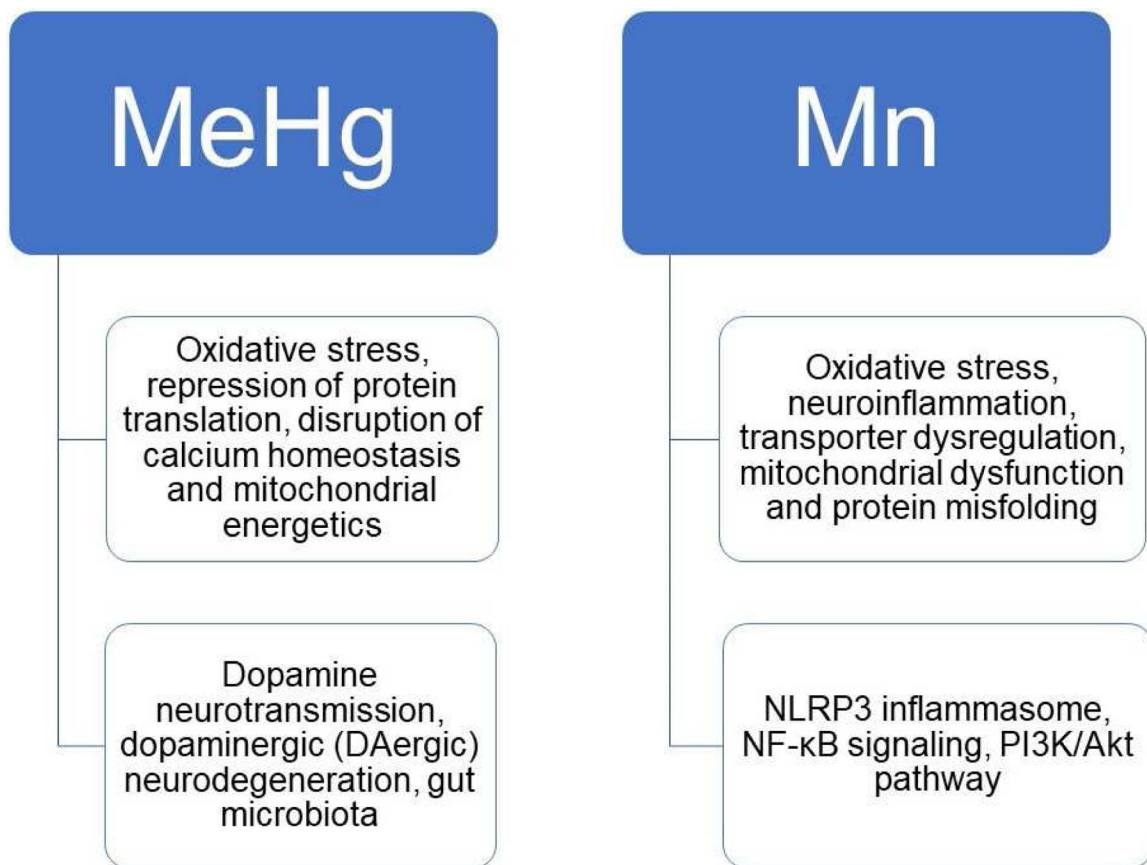


Figure 1:
An overview of the classic and new mechanisms underlying MeHg and Mn induced toxicity.