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## Evaluating a Gene-Environment Interaction in Amyotrophic Lateral Sclerosis: Methylmercury Exposure and Mutated SOD1

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### Abstract

**Purpose of review**—Gene-environment (G×E) interactions likely contribute to numerous diseases, but are often difficult to model in the laboratory. Such interactions have been widely hypothesized for Amyotrophic Lateral Sclerosis (ALS); recent controlled laboratory studies are discussed here and hypotheses related to possible mechanisms of action are offered. Using methylmercury exposure and mutated SOD1 to model the impacts of such an interaction, we interpret evidence about their respective mechanisms of toxicity to interrogate the possibility of additive (or synergistic) effects when combined.

**Recent work**—Recent work has converged on mechanisms of calcium-mediated glutamateexcitotoxicity as a likely contributor in one model of a gene-environment interaction affecting the onset and progression of ALS-like phenotype.

**Summary**—The current experimental literature on mechanisms of metal-induced neuronal injury, and their relevant interactions with genetic contributions in ALS is sparse, but we describe those studies here and offer several integrative hypotheses about the likely mechanisms involved.

#### Keywords

Amyotrophic Lateral Sclerosis; gene-environment (G×E) interaction; methylmercury; AMPA receptor; glutamate; calcium homeostasis

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Conflict of Interest

Jordan M. Bailey, Alexandra Colón-Rodríguez, and William D. Atchison declare that they have no conflict of interest.

Human and Animal Rights and Informed Consent

All reported studies/experiments with animal subjects performed by the authors have been previously published and complied with all applicable ethical standards (including the Helsinki declaration and its amendments, institutional/national research committee standards, and international/national/institutional guidelines).

#### Introduction

Many diseases are thought to arise, or hasten in progression, when environmental events interact with genetic predispositions in unfortunate ways. The idea that gene by environment (G×E) interactions cause disease and dysfunction has been applied to many areas of human health, often successfully expanding our understanding of the complexity of factors that influence health and disease states. Epidemiological reports often herald laboratory studies of disease pathogenesis, but for those thought to arise via  $G \times E$  interactions that path is uniquely complex. Often prototypical environmental events (e.g. exposure to an environmental toxicant and genetic mutations that are known to cause dysfunction) are used to model such an interaction in the laboratory. These studies model a slice of the complexities inherent in the human condition and can yield valuable information regarding the potential mechanisms involved in disease onset and progression, pointing towards meaningful considerations for populations at risk. This review takes a narrow focus on the idea of  $G \times E$  interactions by confining the discussion to a model of one motor neuron (MN) disease, Amyotrophic Lateral Sclerosis (ALS).

No clear genetic (or environmental) factors are known to cause the vast majority (~90%) of ALS cases, which are referred to collectively as sporadic ALS (sALS), making a  $G \times E$ interaction exceedingly likely. Because of this, and because there have been several "outbreaks" of ALS or ALS-like syndromes among non-relatives living near one another, much has been written about environmental risk factors for sALS [1, 2]. Among the most commonly described environmental risks are exposure to pesticides, solvents, and the heavy metals lead and mercury; a history of electric shock or physical trauma; military service; and chronic strenuous physical activity [1]. Some hints at possible epigenetic or genetic predispositions to sALS are also beginning to emerge; carriage of some genetic polymorphisms (e.g. coproporphyrinogen oxidase 4 (CPOX4) [3]; Val66Met in brainderived neurotrophic factor (BDNF) [4]) in combination with mercury exposure was shown to result in heightened motor deficits in humans. Several gene mutations are known to contribute directly to ALS - mutated superoxide dismutase 1 (SOD1), TAR DNA-binding protein 43 (TDP-43), fused in sarcoma/translocated in sarcoma (FUS), and chromosome 9 open reading frame 72 (C9orf72) genes account for around 65% of all familial ALS (fALS) cases and around 11% of all sALS cases [5-7]. Several reviews have examined the possibility that G×E interactions are indeed relevant for ALS and we direct readers to those publications [8–10], as well as to an alternate perspective (see [11]).

The present review assumes a  $G \times E$  interaction *is* relevant to at least some cases of ALS, and instead focuses on the potential neural mechanisms that are affected when genes and the environment converge, manifesting in this disease. While a wealth of circumstantial evidence points toward  $G \times E$  interactions in ALS, studies (experimental or epidemiological) that directly test this hypothesis are uncommon, underpowered or correlational [12]. The few exceptions to this are well-controlled laboratory animal models that permit cause-effect relations to be identified; those using the environmental neurotoxicant methylmercury (MeHg) and a mutated SOD1 gene as exemplar  $G \times E$  factors are emphasized below. We do not propose that only one environmental factor will be important in understanding the complex etiology of ALS, nor do we propose that any environmental factor *alone* will cause

ALS to manifest. We also emphasize that many, divergent genetic predispositions could contribute to this interaction. Indeed, we hypothesize that some individuals might possess currently unrecognized single nucleotide polymorphisms (SNPs) that predispose them to developing ALS when confronted with an environmental insult. Finally, we recognize that more than one cell type inevitably participates in the etiology of ALS – certainly including microglia and astrocytes and likely other neurons that synapse on MNs as well. Much in this field simply remains unknown. Here we discuss the progress that has been made towards untangling the complexities inherent in ALS onset and progression, within the context of the  $G \times E$  interaction model.

#### Amyotrophic Lateral Sclerosis (ALS)

ALS is a neurodegenerative MN disease in which progressive loss of  $\alpha$  and  $\gamma$  MNs of the cerebral cortex, brainstem, and spinal cord occurs [13, 14]. This degeneration first leads to weakness and/or difficulty engaging in voluntary movements. Often symptoms of ALS (e.g. muscle weakness) begin within a single limb or body region [15]. When symptoms begin with speaking or swallowing difficulties it is referred to as bulbar-onset (reflecting the degeneration of MNs in the corticobulbar region of the brainstem). In most cases, partial or total paralysis occurs and respiratory failure or cardiac arrest ultimately results in death. Typically, the onset of ALS is in middle- or late-life, but once it appears its progression is swift; it is generally fatal within 4–5 years following the onset of symptoms [15]. There is no known cure or effective treatment. A number of pharmacotherapies have been tested for the treatment of ALS, however, none to-date have dramatically altered the disease progression. As such, understanding what environmental and/or genetic factors are responsible for the onset and progression of ALS is a goal with significant implications for human health.

Despite the seemingly divergent etiology of sALS and fALS, the clinical manifestation and time course of these forms are nearly identical, suggesting a shared pathophysiology. It is possible that with a disease that is associated with some degree of variability (e.g. in age of onset, progression and survival time) that multiple pathways might be differentially responsible. Some of the most common characteristics attributed to ALS include a high sensitivity of MNs to glutamate (Glu)-mediated damage, difficulty buffering changes to internal Ca<sup>2+</sup> concentration ( $[Ca^{2+}]_i$ ), dysfunction of mitochondrial processes, the generation of reactive oxygen species (ROS), and various alterations to the cytoskeleton [16, 17]. Because ALS emerges relatively late in life, perhaps following a long silent latency, identifying primary (vs. secondary) pathways is made more difficult. Further complicating the issue, emerging evidence suggests that ALS is not a MN-only disease, as once thought; astrocytes and microglia also seem to contribute to its pathophysiology [18, 19].

## Cu<sup>2+</sup>/Zn<sup>2+</sup> Superoxide Dismutase 1 (SOD1) Gene Mutations

Almost all of the mechanisms associated with ALS (mentioned above) have been identified using rodent models, especially mice expressing mutations in the human SOD1, referred to below as SOD1<sup>G93A</sup>. SOD1<sup>G93A</sup> has an amino acid substitution of glycine to alanine at residue 93. This mutation leads to a toxic gain of function of the enzyme. The nature of this

gain in function is as yet unclear, but is the subject of much investigation [20]. SOD1 is a  $Cu^{2+}/Zn^{2+}$  superoxide dismutase that is ubiquitously expressed in all cells. Over 150 mutations have been identified in the SOD1 sequence (http://alsod.iop.kcl.ac.uk) and unsurprisingly mutant SOD1 appears to elicit neurotoxicity via many mechanisms (see [21] for an excellent review on the mechanisms of mutant SOD1 neurotoxicity). The consequences associated with mutant SOD1 involve MNs, microglia, astrocytes and oligodendrocytes and include endoplasmic reticulum (ER) stress, mitochondrial dysfunction, excitotoxicity, oxidative stress, non-cell autonomous toxicity of neuroglia, axonal transport disruption. SOD1 mutations show a characteristic prion-like propagation. It is even possible that *wild-type* SOD1 can become misfolded and ultimately contribute to the pathogenesis of sALS [22, 23]. Some of these mechanisms closely resemble those involved in MeHg's neurotoxicity.

## Methylmercury (MeHg)

The scientific community learned of MeHg's toxicity following a number of catastrophic poisoning events in which discrete populations were exposed to relatively high concentrations of MeHg [24–26]. Perhaps the most infamous were the events in Iraq and Japan occurring in the last century, which have been discussed at length elsewhere [25, 26]. Chronic, low-level exposure to MeHg, however, is of concern for much of the world's population because the consumption of contaminated seafood serves as a primary route of exposure to this toxicant - meals of tuna and swordfish, for example, both deliver relatively high concentrations of MeHg to the consumer and are common worldwide. Some populations for which seafood constitutes a large portion of the diet, or where MeHg contamination is particularly elevated, may be at even greater risk. These include the Amazon rainforest, the Faroe Islands, artisanal gold mining communities, the Great Lakes region of the U.S., Greenland, and perhaps the Seychelles Islands [27–31].

MeHg consumption has been associated with ALS-like syndromes, which has led to its inclusion in animal models examining a G×E interaction in ALS. The characteristic motor impairments following post-birth MeHg intoxication (e.g. exposure to high concentrations) have largely driven its inclusion in the G×E studies highlighted here. Ataxia, visual disturbances, muscle weakness and sensory abnormalities are all common symptoms of MeHg intoxication [26, 32]. These symptoms are often preceded by lengthy asymptomatic periods, generating the characteristic "silent latency" associated with MeHg exposure [33]. Fetal exposure to high concentrations of MeHg presents differently and is associated with physical deformations and cognitive impairment. Notably, unusually high rates of motor disease and dysfunction were observed following the Iraq poisoning episode, in which a large population consumed very high concentrations of MeHg over a relatively short period of time. An abnormally high percent (14%) of this population presented with "myasthenia gravis-like" syndromes [25], a term used loosely to capture the profound motor disturbances that occurred among those exposed. Similarly, ALS-like diseases have emerged following accidental or occupational exposures to high doses of mercurials [34, 35]. Perhaps lower concentrations, like those found in seafood-rich diets could interact with (currently unrecognized) genetic polymorphisms to hasten (or even trigger) the onset of ALS.

Studies of the pathology of MeHg toxicity that use rodent models have demonstrated that 1) MeHg accumulates in, and degenerates, a MNs in the spinal cord, 2) when MNs become saturated with MeHg, ataxia is observed and 3) neurophagia, atrophy and degeneration occur in the anterior horn of the lumbar region of the spinal cord [36-38]. More recent studies have demonstrated that MeHg exposure leads to a concentration-dependent incidence of cell death in MNs that could occur as a result of alterations in intracellular  $Ca^{2+}$  ([Ca<sup>2+</sup>];) homeostasis [39, 40]. Acute *in vitro* exposure to low concentrations  $(0.1 - 1.5 \,\mu\text{M})$  of MeHg in primary spinal cord MNs leads to alterations in  $[Ca^{2+}]_i$ . Disruption of  $[Ca^{2+}]_i$  is mediated in part by the ionotropic Glu receptors: N-methyl-D-aspartate (NMDA) and a-amino-3hydroxy-5-methyl-4-isoxazolepropionic acid (AMPA) [40]. Taken together, these studies demonstrate that MNs degenerate after exposure to MeHg, and that this can occur as a result of the observed alterations in  $[Ca^{2+}]$ ; that are mediated by ionotropic Glu receptors. This observation is similar to one mechanism of pathogenesis observed in ALS. Studies in other neuronal cells, including cerebellar granule cells, have shown that MeHg exposure causes alterations in neurotransmitter release resulting in an increase of Glu release [41] and  $[Ca^{2+}]_i$ [40, 42], mitochondrial damage [43], increases in ROS [44, 45], and impaired function of the excitatory astrocytic Glu transporter (excitatory amino acid transporter 2 (EAAT2)) [46]. Again, these are among the mechanisms of toxicity observed in sALS and fALS [13].

Clearly, MeHg exposure is associated with a number of deleterious effects on cells, but the ability for MeHg to disrupt divalent cation regulation (and the related excitotoxicity that occurs as a result) is of particular interest with respect to ALS [40]. Even very low concentrations of MeHg (e.g. sub- and low- $\mu$ M) can increase levels of  $[Ca^{2+}]_i$  in single cells or synapses [47–49] and precipitate cell death following a delay. The spontaneous release of Glu is stimulated by MeHg, which can itself disrupt the homeostasis of intracellular divalent cations [41]. MeHg's effects on mitochondrial Ca<sup>2+</sup> have been shown to contribute to elevated levels of  $[Ca^{2+}]_i$  and cell death [43, 50]. Behavioral deficits, including both motoric and cognitive dysfunction, that result from MeHg exposure have been linked to disrupted  $[Ca^{2+}]_i$  homeostasis [51]. Further, the oral administration of a Ca<sup>2+</sup> channel antagonist concomitant with MeHg exposure ameliorated the behavioral toxicity associated with MeHg [51] in rodents. Thus,  $[Ca^{2+}]_i$  homeostasis disruption is also a valid mechanism at the level of the whole organism.

#### Evidence of a G×E Interaction

Recently, the explicit focus of our own work has been to determine if a G×E interaction of metal-induced neuronal injury can contribute to the development (or progression) of ALS. We have successfully demonstrated that the time course of mutant SOD1<sup>G93A</sup>-induced motor dysfunction is hastened by chronic MeHg exposure [52]. Mice expressing the human SOD1<sup>G93A</sup> mutation were given chronic, low concentrations of MeHg as they aged. These "dual-hit" animals displayed profound motor dysfunction (e.g. rotorod failure) weeks before the unexposed SOD1 mutants; wild type animals receiving MeHg at the same level *never* displayed this motor impairment. At the time of impairment, these animals had elevated  $[Ca^{2+}]_i$  in MNs of the brainstem. Death and dysfunction of brainstem MNs models "bulbar onset" ALS – this occurs when symptoms begin in the craniofacial regions. This excess  $[Ca^{2+}]_i$  was associated with  $Ca^{2+}$ -permeable AMPA receptors: the AMPA/Kainate receptor

antagonist 6-cyano-7-nitroquinoxaline-2,3-dione (CNQX) delayed the MeHg-induced increase in  $[Ca^{2+}]_i$  in all SOD1<sup>G93A</sup> mice (irrespective of MeHg exposure) but the Ca<sup>2+</sup> permeable AMPA receptor antagonist 1-naphthyl acetyl spermine (NAS) further reduced  $[Ca^{2+}]_i$  in the MeHg exposed SOD1 mutants compared to those untreated with MeHg [52]. These results support a hypothesis that a MeHg-induced increase in Ca<sup>2+</sup> and cell death of MNs are mediated in part by Ca<sup>2+</sup> permeable AMPA receptors, which enhances the dysfunction associated with the SOD1 mutation.

Increases in [Ca<sup>2+</sup>]<sub>i</sub> among these "dual hit" animals was replicated in spinal cord MNs [53], under conditions of acute MeHg exposure (in spinal cord tissue). The spinal cord is a target of particular interest in understanding ALS etiology, as it models the limb onset (or, nonbulbar) forms of this disease. Intracellular Ca<sup>2+</sup> regulation was tracked via fluo-4 epifluorescence following acute 20µM MeHg exposure in spinal cord slice from SOD1G93A mutants and their appropriate controls. As a function of exposure time, acute treatment with MeHg caused a greater increase in fluo-4 fluorescence in the spinal cord tissue of SOD1<sup>G93A</sup> mice compared to control mice. These data further suggest that MeHg interacts with the SOD1<sup>G93A</sup> mutation to enhance neuronal dysfunction (mediated via intracellular Ca<sup>2+</sup> dysregulation among neurons of the spinal cord). This, again, likely demonstrates evidence of a G×E interaction relevant to the etiology of ALS. In both cases, an environmental event was shown to speed the onset and progression of the ALS phenotype caused by the SOD1 mutation (i.e. the genetic contribution) in a way that was not seen with either the gene or environmental event alone. Ongoing and future projects include other genetic models of ALS and as additional genetic targets become known this paradigm can be expanded and tested within numerous (genetic) contexts. These data are of conceptual importance as they demonstrate, unequivocally, that the *possibility* of a G×E interaction in the onset and progression of this disease exists.

Understanding the mechanism(s) underlying this interaction should prove useful for identifying the potential targets of *other* environmental and genetic factors relevant to ALS. In brief, we have hypothesized ([based partly on [52]) that exposure to MeHg disrupts MN divalent cation homeostasis, thereby stimulating MN Glu release. MeHg subsequently inhibits astrocyte Glu uptake by excitatory amino acid transporters (EAATs) [54]. In so doing, MeHg affects the functional relationship between MNs and their attendant astrocytes and generates ROS [54]. However, the proximal events in this process could be MN- or astrocyte-directed. So by using MeHg-treated SOD1 mice with distinct SOD1 mutations we have begun to address the temporal relationship that exists between astrocyte and MN-directed effects. The interdependence of this pair may ultimately contribute to an enhanced development of ALS phenotype. The concomitant effects of increased ROS generation and Ca<sup>2+</sup>-dependent excitotoxicity are likely accelerating the course of ALS development.

#### Convergence on a Mechanism: Glutamate-Induced Excitotoxicity

 $Ca^{2+}$ -mediated, Glu-induced excitotoxicity is thought to be an important cellular-level mechanism that underlies MN loss in ALS and dysfunction in MeHg toxicity [41] – and by extension, in the G×E interaction discussed here. Increased Glu release results in increased  $[Ca^{2+}]_i$  and activating proteins, which can cause mitochondrial toxicity and the generation of

ROS. Moreover, because both MeHg and ALS are associated with increased ROS generation, the MeHg-induced increase of oxidative stress likely contributes to an early onset of ALS. An enhancement in the sensitivity of MNs to Glu-induced excitotoxicity [55] is also hypothesized to contribute to the development of ALS, and is the process that might be especially relevant to the increases in Glu release that are associated with exposure to MeHg [41]. Elevated concentrations of Glu and aspartate have been found in the cerebrospinal fluid of patients with ALS [56], which supports this emphasis on Glu-induced excitotoxicity. In this way, excitotoxicity ultimately results in cell damage and death via  $Ca^{2+}$ -mediated pathways. Increased  $[Ca^{2+}]_i$  is integral in this pathogenesis, and recent reports from our laboratory have shown that MeHg exposure can accelerate increases in  $[Ca^{2+}]_i$  in MNs from the brainstem [52] and spinal cord [53] of SOD1<sup>G93A</sup> mutants.

#### The importance of AMPA receptors

The extension of these findings has led to subsequent, ongoing projects by our group, designed to identify the role of Glu receptors in MeHg-induced toxicity. Examining AMPA receptor function has been central to this aim. AMPA receptors are the ligand-gated Glu receptors in the central nervous system that mediate fast excitatory neurotransmission. When one of the four AMPA receptor subunits, GluA2, is not present (or is in an RNA unedited state), AMPA receptors become Ca<sup>2+</sup> permeable. In ALS, increased levels of unedited GluA2 have been observed. Adenosine deaminase acting on RNA 2 (ADAR2) edits GluA2, and in vitro studies have shown that Glu-induced excitotoxicity can cause dysfunction of ADAR2, leading to increased expression of Ca<sup>2+</sup>-permeable GluA2 in neuronal cells [57, 58]. MeHg exposure causes increased Glu release [41], impaired function of excitatory amino acid transporter EAAT2 [46], increased expression of GluA2 in the brainstem [59] and increases in  $[Ca^{2+}]_i$  in MNs mediated through AMPA receptors [40, 52]. One hypothesis here is that chronic MeHg exposure impairs ADAR2 function, leading to an increase in Ca<sup>2+</sup> permeable AMPA receptors and as a result increased [Ca<sup>2+</sup>]<sub>i</sub>. Preliminary evidence has indicated that MeHg exposure increases the expression of AMPA receptor subunits, especially GluA2, and decreases the RNA editing enzyme ADAR2 in brainstem hypoglossal tissue from rats, a region rich in MNs, to a greater extent in SOD1<sup>G93A</sup> MNs compared to control [59]. See Fig.1 for a schematic representation of the mechanisms thought to be relevant to both ALS and MeHg exposure.

Finally, preliminary studies from our group have demonstrated that low-dose MeHg exposure leads to a concentration-dependent cell death following increases in  $[Ca^{2+}]_i$  that are mediated by AMPA receptors and alterations in AMPA receptor expression in human MNs [60]. Thus, if this is the case in normal human MNs, in a population that has a predisposition to ALS due to genetic mutations, degeneration of these cells may occur faster when exposed to MeHg. Based off these preliminary results and supporting literature it is hypothesized that MeHg action in MNs can have an additive (or perhaps synergistic) effect in ALS-predisposed populations – and, this could have been the case in the mouse study [52] that used the SOD1<sup>G93A</sup> mouse model.

#### Conclusions

The result of recent work performed in our lab suggests that Glu-mediated, Ca<sup>2+-</sup>dependent MN damage can result from a G×E interaction. This was among the first definitive demonstrations of a specific contribution of a persistent environmental contaminant to the development of the ALS phenotype in animal models. MeHg induces multiple effects on neurons that could heighten their sensitivity to subsequent Glu-mediated excitotoxicity due to disease process, SNPs or additional exposure to environmental agents. It also disrupts Glu uptake by astrocytes, which would exacerbate such effects.

The underlying hypothesis here, on which the work reviewed above is based, is that some individuals might possess currently unrecognized SNPs that predispose them to developing ALS when confronted with an environmental insult. In this way, an environmental insult might act to tip the balance of cell damage in the favor of this disease (and possibly others). Sufficient evidence already exists linking exposure to many environmental chemicals including metals and pesticides, which are ubiquitous in the environment, to MN disease. Considering the vast prevalence of diseases that lack any clear genetic cause and the known impact of one's environment or lifestyle on health outcomes, it is prudent to examine seriously the role of environmental contributions to such diseases. The scope of the present review was narrowed to include an examination of one model of the G×E phenomenon: the impact of MeHg exposure on the onset and progression of a "humanized" mouse expressing an SOD1 mutation model of ALS. It is worth noting that it is not the intention of the present review to propose that MeHg alone *causes* ALS and it is acknowledged that some of the genetic predispositions to ALS are at present unknown. What is apparent is that a further understanding of these genetic and environmental risk factors, and how they interact, will be integral to a prevention of disease via the reduction of exposures to environmental contributors, particularly for potentially susceptible individuals.

One important implication of the glutamate-induced excitotoxicity that we discuss here, within the context of MeHg exposure, is that other environmental factors that similarly disrupt calcium homeostasis or otherwise elicit a similar trajectory of neuronal dysfunction should be carefully considered as relevant to ALS disease progression. We would also like to acknowledge that, although limited to-date, evidence is continuing to mount that demonstrates the relevance of  $G \times E$  interactions in ALS (beyond MeHg exposure). For instance, the administration of statins (HMG-CoA reductase inhibitors) has been shown to hasten the ALS-like phenotype of SOD1 mutant rodents [61]; exposure to the toxin Betamethylamino-L-alanine (BMAA) hastens motor dysfunction in zebrafish SOD1 mutants [62]; and, finally, there is emerging interest in examining  $G \times E$  interactions that are protective against the ALS-like phenotype seen in mouse models of ALS (e.g. dietary restriction [63], vitamin D3 supplementation [64]), which, in a different way also addresses the interactions among environment and genetics in the onset and progression of ALS.

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## Abbreviations

G×E	Gene, environment interaction
ALS	Amyotrophic Lateral Sclerosis
sALS	sporadic Amyotrophic Lateral Sclerosis
fALS	familial Amyotrophic Lateral Sclerosis
SOD1	superoxide dismutase 1
FUS	fused in sarcoma/translocated in sarcoma
TDP-43	TAR DNA-binding protein 43
C9orf72	chromosome 9 open reading frame 72
MN	motor neuron
CPOX4	coproporphyrinogen oxidase 4
BDNF	brain-derived neurotrophic factor
MeHg	Methylmercury
SNP	single nucleotide polymorphisms
Glu	glutamate
[Ca <sup>2+</sup> ] <sub>i</sub>	internal calcium concentration
NMDA	N-methyl-D-aspartate
AMPA	a-amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid
ROS	reactive oxygen species
EAAT	excitatory amino acid transporter
CNQX	6-cyano-7-nitroquinoxaline-2,3-dione
NAS	1-naphthyl acetyl spermine
GluA2	AMPA receptor subunit 2
ADAR2	adenosine deaminase acting on RNA

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- •Of importance
- ••Of outstanding importance

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#### Figure 1.

Role of AMPA receptors in ALS and MeHg toxicity on MNs. In ALS, AMPA receptors contribute to the alterations in intracellular  $Ca^{2+}$ . This effect is in part due to the decrease in RNA editing of the GluA2, which results from a decrease in ADAR2. That contributes the increase in the unedited form of GluA2 (GluA2Q) containing receptors in MNs. This is one of the mechanisms that contribute to MN cell death in ALS. MeHg-induced  $Ca^{2+}$  dysregulation in MNs is also mediated in part by  $Ca^{2+}$  permeable AMPA receptors. Our preliminary studies have shown that there is a decrease of ADAR2. However, we do not know how and if the GluA2 subunit is affected by MeHg exposure. This should be the focus of future studies.