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Methionine sulfoxide reductase A affects β -amyloid solubility and mitochondrial function in a mouse model of Alzheimer's disease

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Moskovitz J, Du F, Bowman CF, Yan SS. Methionine sulfoxide reductase A affects β -amyloid solubility and mitochondrial function in a mouse model of Alzheimer's disease. *Am J Physiol Endocrinol Metab* 310: E388–E393, 2016. First published January 19, 2016; doi:10.1152/ajpendo.00453.2015.—Accumulation of oxidized proteins, and especially β -amyloid (A β), is thought to be one of the common causes of Alzheimer's disease (AD). The current studies determine the effect of an in vivo methionine sulfoxidation of A β through ablation of the methionine sulfoxide reductase A (MsrA) in a mouse model of AD, a mouse that overexpresses amyloid precursor protein (APP) and A β in neurons. Lack of MsrA fosters the formation of methionine sulfoxide in proteins, and thus its ablation in the AD-mouse model will increase the formation of methionine sulfoxide in A β . Indeed, the novel MsrA-deficient APP mice (*APP⁺/MsrAKO*) exhibited higher levels of soluble A β in brain compared with *APP⁺* mice. Furthermore, mitochondrial respiration and the activity of cytochrome *c* oxidase were compromised in the *APP⁺/MsrAKO* compared with control mice. These results suggest that lower MsrA activity modifies A β solubility properties and causes mitochondrial dysfunction, and augmenting its activity may be beneficial in delaying AD progression.

posttranslational modification; Alzheimer's disease; mitochondria; oxidative stress

OXIDATIVE STRESS OCCURS IN BIOLOGICAL SYSTEMS when generation of reactive oxygen species and reactive nitrogen species exceeds the system's capacity to eliminate these species. Oxidative stress is a major deleterious mechanism in Alzheimer's disease (AD) (8, 35, 42), other neurodegenerative diseases (11), and normal aging (43). In AD, oxidative damage markers, including lipid peroxidation and nitration, nucleic acid oxidation, and protein carbonylation, are increased in vulnerable brain areas relative to age-matched healthy individuals (26).

AD is characterized pathologically by extracellular amyloid plaques comprised predominantly of fibrillar β -amyloid (A β) peptide and intracellular neurofibrillary tangles made of hyperphosphorylated tau (38). Amyloid plaques are surrounded by inflammation, including activated microglia and astrocytes, which contribute to creation and maintenance of oxidative stress (13). Although historically amyloid plaques were thought to cause AD (36), current evidence indicates that the pathological process leading to AD begins with synaptic injury by neurotoxic A β oligomers, whereas formation of plaques and tangles are downstream events (36). Oxidative stress is one of the earliest consequences of toxic insults mediated by soluble

A β oligomers (27). Mitochondria are particularly sensitive to oxidative stress, and reduced metabolic activity resulting from oxidative damage to vital mitochondrial components has been demonstrated in AD (14). Consequently, antioxidant therapy has been associated with reduced risk for AD (7, 35).

Methionine (Met) is highly susceptible to oxidation in vivo, particularly under conditions of oxidative stress. For example, the sulfoxide form has been found to comprise 10–50% of A β in amyloid plaques of AD brain (2, 6, 16, 25), although it is difficult to determine whether its existence in the plaques contributes to AD etiology or results from the highly oxidative environment around the amyloid plaques, where A β may be trapped for long periods. Oxidation of Met to Met(O) is reversible, and the reverse reaction is catalyzed in vivo by the methionine sulfoxide reductase (Msr) system, comprising peptide-methionine (S)-S-oxide reductase (MsrA) and peptide-methionine (R)-S-oxide reductase (MsrB), which reduce the S and R enantiomers, respectively, of the sulfoxide group. Thus, these enzymes provide both an efficient repair mechanism for oxidative damage to Met residues and general protection against oxidative stress by scavenging reactive oxygen species through the recycling of Met (28).

Mammalian MsrA is encoded by a single gene (19) and is found in both the cytosol and mitochondria (12). Studies in *Msr*-knockout (*MsrAKO*) mice have shown increased vulnerability to oxidative stress (21) and oxidative pathology similar to that associated with AD (32) and Parkinson's disease (PD) (29). Conversely, overexpression of MsrA in various organisms has been shown to provide enhanced protection against oxidative stress and extend survival rate (5, 18, 20).

A cell culture study relevant to AD showed elevated MsrA activity and mRNA levels in human neuroblastoma (IMR-32) cells in response to treatment with the sulfoxide form of A β 42, suggesting that the cells sensed the presence of Met(O) in A β and upregulated MsrA to provide enhanced cellular protection (17). Recently, we reported similar findings in primary rat hippocampal and cortical neurons, in which we observed increased total Msr activity ascribed to both MsrA and MsrB in correlation with protection from cell death induced by the sulfoxide forms of A β 40 or A β 42 (22). Furthermore, exposure of wild-type (WT) and *MsrAKO* mouse cortical neurons to A β 42 and Met(O)-A β showed that lack of MsrA abolishes the protective effect induced by Met(O)-A β (22). A different study demonstrated that overexpression of MsrA in differentiated rat pheochromocytoma (PC-12) cells increased significantly the resistance of the cells to A β 42-induced toxicity (15). These cell culture studies suggest that neurons protect themselves

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against oxidative stress in AD by elevated expression of the Msr system.

Studies using WT rats have shown age-related decline in Msr activity unrelated to disease (33, 40, 44), which correlated with a decrease in the mRNA levels of *MsrA* (10). Overall, the combination of human and animal data suggests that the low Msr levels observed in AD brain (10) result both from aging-related decreased transcription and disease-related translational or posttranslational defects. The decline in Msr activity would be expected to lead to gradual accumulation of protein-Met(O). Thus, it is proposed that both compromised Msr activity and the resulting accumulation of protein-Met(O) [including specifically Met(O)-A β] will increase the levels of soluble-A β while compromising mitochondrial function. Accordingly, to facilitate obtaining relevant data to support our hypothesis, we have created a novel mouse model in which the *MsrA* gene is deleted in an AD mouse model that overexpresses amyloid precursor protein (APP) and A β in neurons.

MATERIALS AND METHODS

Creation of *APP*⁺/*MsrAKO* mice and validation of their high A β levels. The transgenic mice overexpressing an isoform of human Alzheimer's A β precursor protein (APP) exhibit AD-like phenotypes at mature to older ages. These mice produce high levels of A β in the brain following cleavage of the overexpressed human APP protein. The APP transgenic mice we used for our studies are Tg (PDGFB-APPSwInd, or "J20") mice that express a mutant form of human APP bearing both the Swedish (K670N/M671L) and the Indiana (V717F) mutations (*APPSwInd*) and were purchased from Jackson's Laboratory (Sacramento, CA) and will be denoted as *APP*⁺ mice. This AD mouse model is well characterized in terms of mitochondrial function, oxidative stress, A β accumulation, and synaptic function (23). Crossing these *APP*⁺ mice with *MsrAKO* mice will result in a hybrid mouse model that overexpresses APP and lacks the *MsrA* gene. Briefly, female *MsrAKO* mice on a C57BL6/129sv background (21) were back-crossed 10 times into a C57BL6 background and mated with males of PDGFB-APPSwInd on a C57BL6 background (*APP*⁺ mice). Double-heterozygous F1s were selected by PCR typing for the human *APP* gene and intercrossed with *MsrA*^{+/-} from F1 to produce F2 progenies that included mice that are homozygous for the *MsrA*^{-/-} allele in combination with heterozygous *APP*⁺ transgenic gene (*MsrAKO*/PDGFB-APPSwInd, named shortly as "*APP*⁺/*MsrAKO* mice"). Littermates that were *MsrA*^{+/-}/PDGFB-APPSwInd served as the *APP*⁺ control mice for the *APP*⁺/*MsrAKO* cohorts. Littermates that were non-*APP*⁺ transgenic *MsrA*^{+/-} mice served as the WT control for the non-*APP*⁺/*MsrAKO* cohorts (*MsrA*^{-/-} mice). All mouse-related experimental protocols were submitted to and approved by the Animal Care Unit Committee of the University of Kansas.

Brain tissue processing. The following indicated brain extractions were performed based on a previously published method (39). Briefly, AD-relevant brain regions (a mixture of hippocampus and cortex) were dissected out of one hemisphere of all mouse brains at euthanasia ($n = 5$ /mouse strain, 7 mo of age). Ice-cold Tris-buffered saline (TBS) consisting of 20 mM Tris-HCl and 150 mM NaCl, pH 7.4, was added to the frozen regions (4:1 vol/wt) in the presence of protease inhibitors mixture (Roche and Life Technologies) and homogenized with a mechanical Dounce homogenizer. The homogenate was spun down at 157,000 g for 15 min at 4°C, and the supernates were aliquoted and stored at -80°C (denoted as the "Tris"-soluble fractions). The pellets were rehomogenized (4:1 vol/wt) in TBS + 1% Triton X-100 and spun as described above. The resultant supernates (denoted as "Triton"-soluble fractions) were aliquoted and stored at -80°C, and the pellet was rehomogenized in TBS + 5 M guanidine HCl. Following the same centrifugation indicated above, the resultant

supernates (denoted as "guanidine"-soluble fractions) were aliquoted and stored as well at -80°C until use. The Tris-soluble fraction contains mainly a mixture of both monomeric and oligomeric A β that is representative of the most soluble species of A β . The quinidine soluble fraction contains mainly aggregative A β species that are representative of the least soluble species of A β . The Triton soluble fraction represents intermediate fraction between the Tris and quinidine soluble fractions that contains mainly a mixture of both oligomeric and aggregative A β species. Having these fractions will facilitate the search for possible correlation between types of A β and their toxicity in brain.

Determination of A β levels. First, to determine that indeed total A β levels were similar in both *APP*⁺ and *APP*⁺/*MsrAKO* brains and higher compared with the non-*APP*⁺ carrier brains (WT and *MsrAKO*), a dot blot analyses were performed. Equal protein amounts of the evenly combined three soluble fractions (Tris, Triton, and guanidine) from each brain sample were applied to a dot blot analyses (in a range of 1–10 μ g protein·dot⁻¹·strain⁻¹). The MOAB-2 antibodies were used as the primary antibody, followed by the secondary antibody horseradish peroxidase (HRP)-conjugated goat anti-mouse IgG (Santa Cruz Biotechnology, Dallas, TX). The MOAB-2 antibodies recognize human as well as mouse A β , unaggregated, oligomeric, and fibrillar forms of synthetic A β 42 and unaggregated A β 40, without recognizing the APP protein (Ref. 45 and personal communication from Dr. Mary Jo LaDu, University of Illinois). The densities of the resulting signals, following exposure of the blot to an X-ray film, were quantified by the National Institutes of Health ImageJ program.

Detection and measurement of Met(O)-A β distribution in brain. We used our anti-Met(O) antibodies to detect Met(O)-A β in the hippocampal and cortex, as it was shown to detect Met(O) in other brain proteins associated with neurodegenerative diseases (3, 30, 31, 41). For this analysis, we have developed a novel sandwich ELISA for the detection of sulfoxidized A β [A β -Met(O)]. The assay worked in the following format: extracted or synthetic A β was captured by an anti-A β antibody (MOAB-2, which is bound to the ELISA 96-well plate), and the rabbit anti-Met(O) antibody is used for the detection of Met(O) of A β . Then, fluorescence visualization is mediated by HRP-conjugated goat anti-rabbit IgG (Santa Cruz Biotechnology). For calibrating the ELISA, synthetic A β (1–40/42) and A β -Met(O) (1–40/42) (a gift from Dr. Gal Bitan, University of California Los Angeles) were used in various concentrations in this ELISA format. The Tris, Triton, and guanidine-soluble fractions of the four mouse strains were loaded into the 96-well plate, and the ELISA assay was performed according to the format described above. Briefly, MOAB-2 antibodies (1:1,000) were added into coating buffer [50 mM carbonate-bicarbonate (pH 9.6)] at ~1.0 μ g/ml, and then they were added to a Nunc-Immuno 96-well plate (Maxisorp Surface Treatment; Thermo-Fisher Scientific, Waltham, MA). The plate was incubated overnight at 4°C and washed with 0.05% (vol/vol) Tween-20 in PBS (PBST). Blocking buffer [0.5% (wt/vol) BSA in PBS] was added to each well and incubated for 1 h at room temperature. Then, the wells were washed with PBST. Equal amounts of proteins from each brain fraction were added to the plate in various dilutions (from 1:20 to 1:100 to eliminate the effect of the Triton-guanidine) and were incubated for overnight at 4°C. The plate was washed with PBST, and anti-Met(O) antibodies were added in PBST (1:1,000) and incubated at room temperature for 2 h. Following washes with PBST, HRP goat anti-rabbit IgG (Santa Cruz Biotechnology) was added to each well (1:5,000) and incubated at room temperature for 2 h. Finally, the plate was washed with PBST, and a mixture of peroxidase substrate solutions was added to each well (Substrate Reagent Pack; R & D Systems-Tocris, Minneapolis, MN). Then, a stopper solution (1 N sulfuric acid) was added, and the plate was read at 450 nm.

Mitochondrial respiration and cytochrome c oxidase expression and activity. Mouse brain regions (a mixture of hippocampus and cortex) were dissected out of one hemisphere of all mouse brains at euthanasia ($n = 5$ /mouse strain). These brain regions were used as a

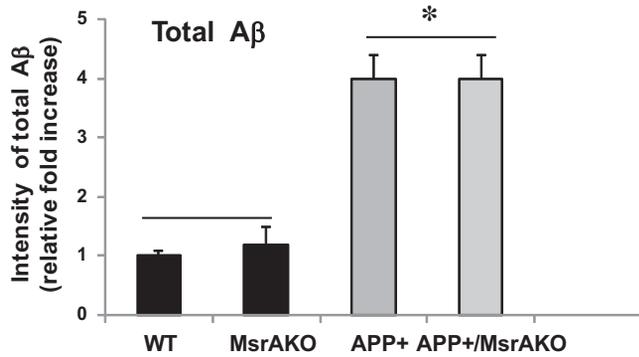


Fig. 1. Levels of total β -amyloid ($A\beta$) in brains of wild-type (WT), peptide-methionine (*S*)-*S*-oxide reductase (*MsrAKO*), amyloid precursor protein (*APP*)⁺, and *APP*⁺/*MsrAKO* mice. Mouse brains (hippocampal and cortical regions) were homogenized and processed as described in MATERIALS AND METHODS. Equal protein amounts of the evenly combined 3 soluble fractions (Tris, Triton, and guanidine) from each brain sample were applied to a dot blot analyses (in a range of 1–10 μ g protein·dot⁻¹·strain⁻¹). The MOAB-2 antibodies were used as the primary antibody, followed by secondary antibody horseradish peroxidase (HRP)-conjugated goat anti-mouse IgG antibodies. The densities of the resulting signals, following exposure of the blot to an X-ray film, were quantified by the NIH ImageJ program. *MsrA*^{-/-} is denoted as *MsrAKO*. *Statistically significant differences between the *APP*⁺ and *APP*⁺/*MsrAKO* group and the WT and *MsrAKO* group ($P < 0.001$).

source for the isolated mitochondrial preparations. Oxygen consumption of each preparation was determined by following mitochondrial respiration assay, using the OROBOROS Oxygraph-2K (OROBOROS Instruments, Innsbruck, Austria), and as described previously (4). The tissue homogenates or isolated mitochondria were resuspended in KCl medium (80 mM KCl, 10 mM Tris-HCl, 3 mM MgCl₂, 1 mM EDTA, and 5 mM potassium phosphate, pH 7.4). Various substrates and inhibitors for mitochondrial respiratory chain complexes were used as described in Figs. 3 and 4. OROBOROS DatLab software was used to calculate the oxygen consumption. The cytochrome *c* oxidase activities of mitochondrial fractions were measured

with a cytochrome *c* oxidase kit (Sigma-Aldrich, St. Louis, MO). Briefly, mitochondrial fraction and enzyme solution were added to the assay buffer. The reaction was initiated by the addition of ferrocyanochrome *c* substrate solution. Enzyme kinetics was determined by following changes in OD values at 550 nm at 10-s intervals (using a Synergy H1 monochromator-microplate reader; Biotek, Winooski, VT). The expression levels of cytochrome *c* oxidase and heat shock protein 60 (HSP60; mitochondrial marker) were determined by a Western blot analysis performed on mitochondrial protein fractions made from the four mouse types (primary antibody: anti cytochrome *c* oxidase subunit VIb and anti-HSP60 antibodies; Abcam, Cambridge, MA).

Statistical analyses. All data values in the presented figures are displayed as means \pm SD. All analyses were run by two-way analysis of variance (ANOVA), followed by Newman-Keuls test for post hoc comparisons.

RESULTS

We have created the *APP*⁺/*MsrAKO* mice as described in MATERIALS AND METHODS. It is worth noting that the rate of success of having *APP*⁺/*MsrAKO* pups was below the expectations. Only 1% of all resulting littermates contained the desired genotype (we have obtained only 5 out of 500 resulting pups from the F2 progenies, as described in MATERIALS AND METHODS). This phenomenon may indicate that complete absence of *MsrA* causes enhanced toxicity of $A\beta$, which negatively affects the survival of the newborn pups. The obtained *APP*⁺/*MsrAKO* pups were grown until they reached ≥ 7 mo of age before any analyses was performed, since the *APP*⁺ mice start to exhibit AD-associated phenotypes only at that age. Accordingly, we have first determined the levels of soluble $A\beta$ in the four mouse strains. As predicted, a significantly high increase in $A\beta$ levels was observed in the brain fractions of the *APP*⁺ carriers compared with the non-*APP*⁺ carrier brains (Fig. 1). Additionally, the levels of total $A\beta$ were compatible in both the *APP*⁺ carrier and the non-*APP*⁺ carrier, suggesting

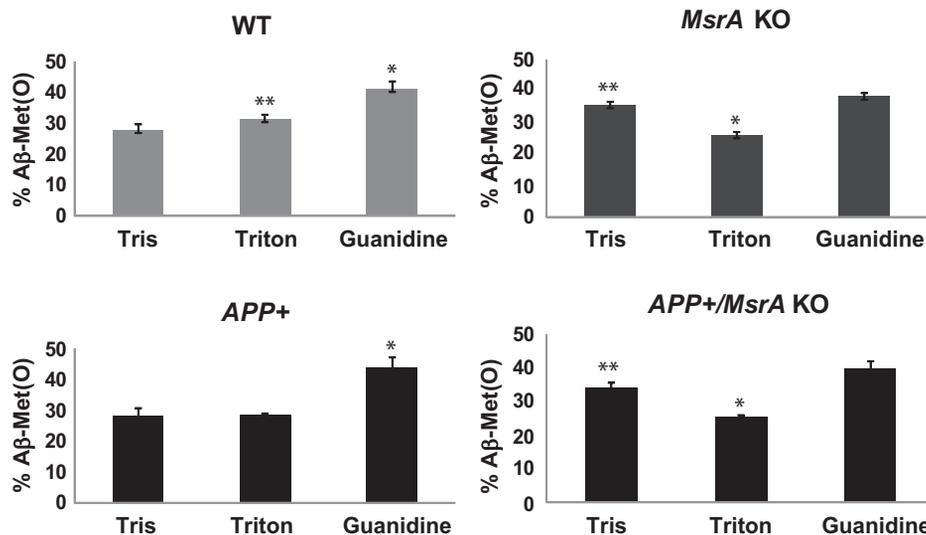


Fig. 2. Levels of $A\beta$ in brains of WT, *MsrAKO*, *APP*⁺, and *APP*⁺/*MsrAKO* mice (7 mo of age). Mouse brains (hippocampal and cortical regions) were homogenized and processed as described in MATERIALS AND METHODS, resulting in Tris-, Triton-, and guanidine-soluble fractions. A sandwich ELISA for the detection of sulfoxidized $A\beta$ [$A\beta$ -Met(O)] was performed using anti- $A\beta$ antibody (MOAB-2) as the capture antibody, rabbit anti-Met(O) antibody for the detection of Met(O) of $A\beta$, and the HRP-conjugated goat anti-rabbit IgG as the signaling probe. *MsrA*^{-/-} is denoted as *MsrAKO*. Statistically significant differences: for WT, * $P < 0.001$ between guanidine and Tris and Triton fractions and ** $P < 0.02$ between Tris and Triton fractions; for *APP*⁺, * $P < 0.001$ between guanidine and Tris and Triton fractions; for *MsrAKO*, * $P < 0.001$ between Triton and Tris and Guanidine fractions and ** $P < 0.04$ between Tris and guanidine fractions; for *APP*⁺/*MsrAKO*, * $P < 0.001$ between Triton and Tris and guanidine fractions and ** $P < 0.01$ between Tris and guanidine fractions.

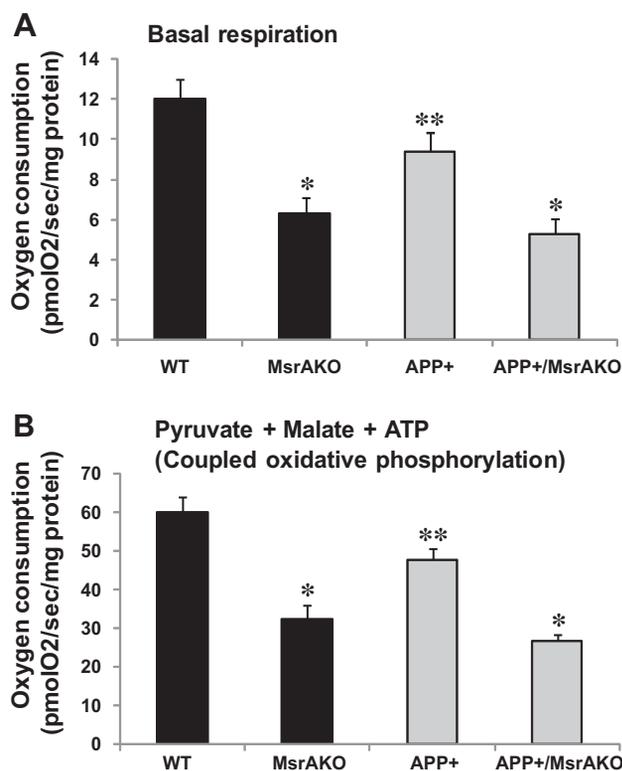


Fig. 3. Mitochondrial respiratory chain activity is impaired in freshly isolated mitochondria in brains of WT, *MsrAKO*, *APP*⁺, and *APP*⁺/*MsrAKO* mice (7 mo of age). Basal respiration (A) and pyruvate + malate + ADP (coupled oxidative phosphorylation) (B). The respiratory values were measured using OROBOROS Oxygraph-2K as described, in MATERIALS AND METHODS. Values are means \pm SD; $n = 5$. * $P < 0.001$ vs. *APP*⁺ and WT groups in A and B; ** $P < 0.05$ vs. WT (1-way ANOVA with Tukey's post hoc comparison).

that ablation of *MsrA* in either WT or *APP*⁺ mice did not have an effect on the expression of A β levels. Then, using our novel ELISA assay, we determined the relative ratio of Met(O) levels in the Tris, Triton, and guanidine-soluble fractions containing mainly the monomeric and oligomeric A β , oligomeric and aggregative A β , and aggregative A β species, respectively. The added signal values obtained from the respective three fractions of each mouse represented its 100% level of A β -Met(O). Accordingly, the distribution of A β -Met(O) levels in all of the fractions in each mouse strain is provided as percent of total A β -Met(O). The ability of the anti-Met(O) antibody to specifically recognize A β -Met(O) was further confirmed by dot blot and our ELISA analyses using synthetic A β and A β -Met(O). As shown in Fig. 2, the pattern of A β -Met(O) distribution exhibited in *MsrA* carriers (WT and *APP*⁺ mouse strains) is similar to and different from the A β -Met(O) distribution exhibited in the *MsrA*^{-/-} carriers (*MsrAKO* and *APP*⁺/*MsrAKO* mouse strains). Compared with the mouse strains having an intact *MsrA* gene, the lack of the *MsrA* gene caused a reduction in the levels of the A β -Met(O) in the guanidine- and Triton-soluble fractions and an increase in the A β -Met(O) levels in the Tris-soluble fraction (Fig. 2). This increased presence of A β -Met(O) in the Tris fraction of the *MsrAKO* and *APP*⁺/*MsrAKO* brain extracts supports the hypothesis that there is a shift from aggregates to the soluble oligomers of MetO-A β in mice lacking *MsrA*. These soluble oligomers are thought to possess more toxic elements to neurons and synapses than

aggregative A β forms alone (37, 39). Furthermore, since *MsrA* is also transported into mitochondria (12), we examined the effect of *MsrA* ablation on mitochondrial function in the presence or absence of A β overexpression. Oxygen consumption of each preparation was determined by following mitochondrial respiration assay, using the OROBOROS Oxygraph-2K. In addition, various substrates and inhibitors for mitochondrial respiratory chain complexes were used as

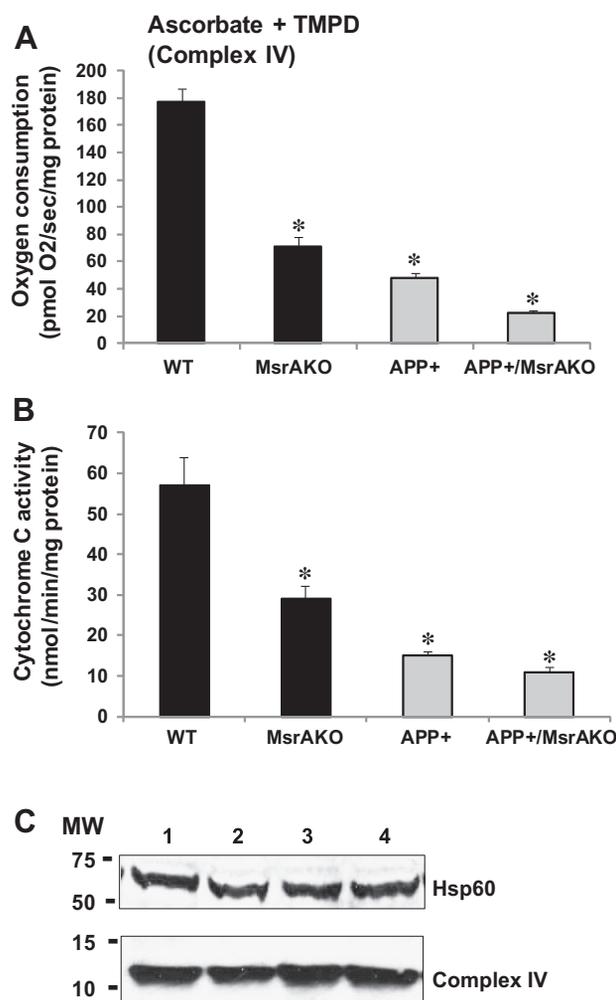


Fig. 4. Cytochrome *c* oxidase (complex IV) mitochondrial function and activity decline in mitochondria of *MsrAKO*, *APP*⁺, and *APP*⁺/*MsrAKO* brains (7 mo of age). A: complex IV mediated oxygen consumption in mitochondria, which was measured using OROBOROS Oxygraph-2K, as described in MATERIALS AND METHODS. Complex IV respiration was calculated as the portion sensitive to potassium cyanide (KCN), a specific inhibitor of cytochrome *c* oxidase (first, the ascorbate-TMPD respiratory rates in the presence and absence of KCN were obtained, and then the final respiratory rate was obtained by subtracting the KCN-insensitive respiration). B: cytochrome *c* oxidase activities of mitochondrial fractions were measured with a cytochrome *c* oxidase kit (Sigma). Briefly, mitochondrial fraction and enzyme solution were added to the assay buffer. The reaction was initiated by the addition of ferrocytochrome *c* substrate solution. Enzyme kinetics was determined by following changes in optical density values at 550 nm at 10-s intervals (using a Synergy H1 monochromator-microplate reader; Biotek). C: protein expression levels of cytochrome *c* oxidase and heat shock protein 60 (HSP60; loading control as a mitochondrial marker) were determined by Western blot analyses (a represented blot is shown, in which each lane contains a pool of equal mitochondrial protein extracts originating from 5 animals/mouse group). Values are means \pm SD; $n = 5$. * $P < 0.001$ vs. WT (1-way ANOVA). MW, molecular weight markers.

described in Figs. 3 and 4. Basal respiration was significantly reduced in the *MsrA* KO mice and *APP*⁺/*MsrA*KO mice compared with WT control and *APP*⁺ control, respectively. The mouse values pattern presented in Fig. 3A was reproduced in the coupled oxidative phosphorylation experiments, as shown in Fig. 3B, with the expected higher oxygen consumption values. Ascorbate and *N,N,N',N'*-tetramethyl-*p*-phenylenediamine (TMPD) were added to the respiratory mixture to determine the capacity of cytochrome *c* oxidase (complex IV). TMPD is an artificial redox mediator that assists the transfer of electrons from ascorbate to cytochrome *c*. Complex IV respiration was calculated as the portion sensitive to potassium cyanide (KCN), a specific inhibitor of cytochrome *c* oxidase (first, the ascorbate-TMPD respiratory rates in the presence and absence of KCN were obtained, and then the final respiratory rate was obtained by subtracting the KCN-insensitive respiration). Respiration rates with ascorbate plus TMPD were decreased by ~60% in the *MsrA*KO compared with WT mice, and a similar reduction was observed in the *APP*⁺/*MsrA*KO compared with *APP*⁺ mice (Fig. 4A). These data show that the lack of MsrA caused a similar negative impact on mitochondrial respiration (mediated by compromising complex IV function) of both WT and *APP*⁺ mice. Furthermore, under inhibitory conditions of complex IV by KCN, the *APP*⁺/*MsrA*KO mice were the most affected. These data strongly suggest that the mitochondrial MsrA activity contributes for maintaining complex IV function, which may be of most importance especially when the mitochondrial function of complex IV in the *APP*⁺ mice is already inhibited (Fig. 4A). Thus, it is suggested that the mitochondrial MsrA's ability to reduce Met(O) residues of either complex IV or other related proteins may contribute to the complex IV-mediated respiration. Indeed, monitoring the enzymatic activity of cytochrome *c* oxidase in these mice showed that the activity paralleled complex IV-dependent respiration changes (Fig. 4, A vs. B). In addition, no significant changes in the expression levels of either cytochrome *c* oxidase HSP60 (a marker of mitochondria) were observed as determined by a Western blot analyses performed on mitochondrial protein fractions made from the four mouse types (Fig. 4C). Accordingly, it is suggested that the observed declined complex IV respiration (Fig. 4A) is associated with Met oxidation of cytochrome *c* oxidase and/or other related mitochondrial proteins to which the *APP*⁺/*MsrA*KO mice are mostly vulnerable.

DISCUSSION

Modification of proteins by methionine oxidation can change their biochemical function and/or structure. Accordingly, the oxidation of the sole methionine residue of A β causes it to lose its fibrillation, and overexpression of *MsrA* leads to a protective effect against neuronal cell death in culture in the presence of either A β or A β -Met(O) (15, 17, 22). In the current studies, we show that lack of MsrA promotes a shift of aggregated forms of A β toward soluble oligomers in vivo (Fig. 2). Since soluble oligomers are thought to possess more toxic elements to neurons and synapses than aggregative A β forms alone (37, 39), it is suggested that enhancing MsrA activity by compounds that can upregulate its transcription may have therapeutic application. For example, we have showed that pergolide, pergolide sulfoxide, and *S*-adenosyl methionine

can upregulate MsrA transcription and activity in cultured neuronal cell (9). Alternatively, immunization of the *APP*⁺ type of mice with methionine sulfoxide-rich protein caused a reduction in amyloid plaque burden in the hippocampus, presumably through clearance of protein-Met(O) [including A β -Met(O)] from brain (22). The role of MsrA in cellular protection against oxidative stress is also evident from its presence in mitochondria, although the regulatory process of its transport into the mitochondria is not clear yet. Evidence for the link between mitochondrial dysfunction and AD has been accumulated in various studies on the role of mitochondria in normal aging and neurodegenerative diseases. For example, reduced cytochrome *c* oxidase activities have been reported in both platelets and brains of mild cognitive impairment and AD patients (1, 24), and this activity reduction was attributed to an enhanced mitochondrial oxidative stress occurring especially at an advanced age. As shown in Figs. 3 and 4, lack of MsrA exacerbates mitochondrial malfunction that is observed in the *APP*⁺ mice. Thus, this situation may mimic the reduction of MsrA expression with age (33, 40, 44), which is suggested to be a risk factor for AD-related mitochondrial malfunction when A β levels are elevated. Further studies are needed to elucidate the interaction between A β -Met(O), mitochondrial function, and neuronal cell death.

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DISCLOSURES

All authors declare that they do not have any potential conflicts of interest, financial or otherwise.

AUTHOR CONTRIBUTIONS

J.M., F.D., and S.Y. conception and design of research; J.M., F.D., and C.F.B. performed experiments; J.M., F.D., C.F.B., and S.Y. analyzed data; J.M., F.D., and S.Y. interpreted results of experiments; J.M. prepared figures; J.M. drafted manuscript; J.M. and S.Y. edited and revised manuscript; J.M. approved final version of manuscript.

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