The protective role of Bax Inhibitor-1 against chronic mild stress through the inhibition of monoamine oxidase A

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Received 13 August 2013
Accepted 15 November 2013
Published 2 December 2013

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The anti-apoptotic protein Bax inhibitor-1 (BI-1) is a regulator of apoptosis linked to endoplasmic reticulum (ER) stress. It has been hypothesized that BI-1 protects against neuron degenerative diseases. In this study, BI-1−/− mice showed increased vulnerability to chronic mild stress accompanied by alterations in the size and morphology of the hippocampi, enhanced ROS accumulation and an ER stress response compared with BI-1+/+ mice. BI-1−/− mice exposed to chronic mild stress showed significant activation of monoamine oxidase A (MAO-A), but not MAO-B, compared with BI-1+/+ mice. To examine the involvement of BI-1 in the Ca2+-sensitive MAO activity, thapsigargin-induced Ca2+ release and MAO activity were analyzed in neuronal cells overexpressing BI-1. The in vitro study showed that BI-1 regulates Ca2+ release and related MAO-A activity. This study indicates an endogenous protective role of BI-1 under conditions of chronic mild stress that is primarily mediated through Ca2+-associated MAO-A regulation.

Bax inhibitor-1 (BI-1) is an evolutionarily conserved, multi-transmembrane protein that is located in the endoplasmic reticulum (ER) and provides cytoprotective functions in both animals and plants. The cytoprotective function of BI-1 was originally discovered in cDNA library screens for human proteins capable of suppressing the death induced by ectopic expression of mammalian Bax protein in yeast. When overexpressed, the anti-apoptotic protein BI-1 has been shown to reduce ER Ca2+ concentrations, possibly through a mechanism involving inositol 1,4,5-trisphosphate receptor (IP3R) activation. Knockdown of BI-1 results in increased ER basal Ca2+ concentration and increased Ca2+ release into the cytosol following thapsigargin challenge. The regulatory role of BI-1 in Ca2+ may contribute to its neuroprotective effects. As such, BI-1 has recently been identified as a critical constituent in both ischemia and traumatic brain injury mouse models, showing regulatory effects against serotonin-depletion stress. Disturbances in intracellular Ca2+ homeostasis have been reported in depression and bipolar diseases. These disruptions hinder cell viability and function. Although BI-1 also confers resilience in animal models, the mechanism by which BI-1 protects against depression is not clearly understood.

Neuronal Ca2+ is linked both directly and indirectly to monoamine oxidases (MAOs), the core enzymes for monoamine metabolism. The MAOs, which are located in the mitochondrial membrane, metabolize monoamines and contribute to the maintenance of monoamine homeostasis. Biochemical and pharmacological studies have revealed two isoforms of MAO, MAO-A and MAO-B. While MAO-A oxidizes serotonin, norepinephrine and dopamine, MAO-B preferentially oxidizes dopamine. Ca2+ accumulation leads to enhanced MAO activity in neuronal cells and neuronal disease models.

In this study, we tested the hypothesis that BI-1 is involved in Ca2+-related psychiatric conditions including depression through endogenous Ca2+ homeostatic regulation. This study linked MAO-A activity with BI-1-associated Ca2+ regulation in a chronic mild stress model.
Results

BI-1 mice are more vulnerable to stress-induced depression-like behavior than BI-1\textsuperscript{+/+} mice. BI-1\textsuperscript{+/+} and BI-1\textsuperscript{−/−} mice aged 7 to 8 months were exposed to chronic mild stress for 2 or 6 weeks according to the experimental design (Supplemental Fig. 1). During the stress procedures, body weight was significantly more decreased in BI-1\textsuperscript{−/−} mice than in BI-1\textsuperscript{+/+} mice (Supplementary Fig. 2). Because corticosteroid is considered to be a stress marker\textsuperscript{16-17}, the level of serum corticosterone was compared between BI-1\textsuperscript{+/+} and BI-1\textsuperscript{−/−} mice with or without stress. In BI-1\textsuperscript{−/−} mice, the corticosteroid concentration was greatly increased under conditions of stress compared with BI-1\textsuperscript{+/+} mice, especially in the 6-weeks group, with the BI-1\textsuperscript{−/−} group showing high sensitivity to stress (Fig. 1A). Sucrose intake, another representative marker for the stress response, was significantly decreased in the 6-weeks stress group of BI-1\textsuperscript{−/−} mice compared with BI-1\textsuperscript{+/+} mice (Fig. 1B). The response to stress was also compared between groups through analysis of locomotor activity. Significant differences were observed in distance traveled and locomotor time between the two groups under the 6-weeks chronic mild stress regime (Fig. 1C, Supplementary Table 1); these two parameters were significantly higher in the BI-1\textsuperscript{−/−} mice relative to the wild-type mice, suggesting that knockout of BI-1 increases vulnerability to chronic mild stress. Forced swimming, which imposes high stress and evokes marked changes in physical and emotional components\textsuperscript{18}, had a similar effect on swimming time in both BI-1\textsuperscript{+/+} and BI-1\textsuperscript{−/−} stressed mice (Supplementary Fig. 3).

BI-1 affects changes in hippocampus volume, ROS accumulation, BDNF expression and ER stress associated with chronic mild stress. Since the hippocampus is known to be vulnerable to stress that induces depression\textsuperscript{19}, the change in the volume of the hippocampus of BI-1\textsuperscript{+/+} and BI-1\textsuperscript{−/−} mice was measured by MRI.
image analysis. Representative MRI images of hippocampi are shown in Figure 2A. Hippocampus volume was quantified and compared between BI-1+/+ and BI-1−/− mice. The volume of the hippocampus in the BI-1 knockout mice was significantly reduced in the 6-weeks stress group (Fig. 2B). In addition, hippocampus structure was analyzed through hematoxylin and eosin (H&E) staining. Although the hippocampal morphology of mice exposed to 2 weeks of stress was similar to that of controls, hippocampus structural alterations were observed after 6 weeks of stress and were more dramatic in the BI-1 knockout mice (Fig. 2C).

Dihydroethidium staining was performed in frozen sections to examine reactive oxygen species (ROS) involvement in the chronic mild stress response. As expected, we observed ROS accumulation over the 2- and 6-weeks stress periods, especially in the BI-1 knockout condition (Fig. 2D, quantified in the lower panel). BI-1+/+ mice showed positive staining for brain-derived neurotrophic factor (BDNF) in the hippocampus, with a decrease in expression during the 6-weeks stress period. Compared with the staining pattern in BI-1+/+ mice, BDNF was not positively identified over the 6-weeks stress condition in BI-1−/− mice, particularly in the dentate gyrus (Fig. 3A). Based on these data, it was necessary to examine the ER stress response, because BI-1 is an established ER stress regulator and ER stress regulation has been suggested as a possible therapeutic mechanism for psychiatric diseases including depression20,21. Indeed, the ER stress response was more clearly observed in the hippocampus following exposure to chronic mild stress in the BI-1−/− mice compared with the BI-1+/+ mice, as demonstrated by immunostaining of representative ER stress response proteins GRP78 and CHOP and cleaved caspase-3 (Fig. 3B). As shown in Figure 3C, GRP78, CHOP, p-IRE-1α, and spliced XBP-1 were clearly expressed in the BI-1−/− mice exposed to 6 weeks of chronic mild stress, consistent with vulnerability to stress in the BI-1 knock-out condition. Taken together, these morphological and signaling changes in the hippocampus suggest that stress mainly affects the hippocampus region, where BI-1 regulates ROS accumulation and ER stress leading to alterations in the structure of the hippocampi.

**BI-1 regulates MAO-A activity, leading to the regulation of serotonin, norepinephrine or dopamine.** ER stress is also associated with the disturbance of ER function including the disruption of Ca2+ homeostasis. Since Ca2+ disturbance increases the activity of monoamine oxidase-A (MAO-A), a mitochondria-bound enzyme that generates ROS as a natural by-product of the deamination of neurotransmitters such as serotonin22, MAO-A and MAO-B enzyme activities were analyzed in brains from BI-1+/+ and BI-1−/− mice following exposure to chronic mild stress. In the BI-1−/− mice, the activity of MAO-A, but not MAO-B, was significantly increased, especially in mice exposed to 6 weeks of chronic mild stress (Fig. 4A). MAO-A and MAO-B mRNA and protein expressions were also analyzed in BI-1+/+ and BI-1−/− mice following exposure to chronic mild stress and the fluorescence intensity was quantified (lower).

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**Figure 2** | BI-1 knock-out enhances hippocampus volume and structure changes and ROS accumulation under chronic mild stress. (A) Brain images and (B) hippocampus volumes from BI-1+/+ and BI-1−/− mice were analyzed after 6 weeks of chronic mild stress. (C) H&E staining of BI-1+/+ and BI-1−/− mouse brains was performed after 2- and 6-weeks periods of chronic mild stress. (D) Dihydroethidium staining of BI-1+/+ and BI-1−/− mouse brains was performed after 2- and 6-weeks chronic mild stress treatment and the fluorescence intensity was quantified (lower).
to chronic mild stress. No differences were found between the two groups of mice (Supplementary Fig. 4A, B). These data suggest that BI-1 affects MAO-A activity post-translationally. The concentrations of representative neurotransmitters were analyzed through LC-MS-MS analysis. Levels of the monoamines serotonin, norepinephrine and dopamine in whole brains from BI-1^2/2^ mice were significantly decreased after 6 weeks of chronic mild stress compared with the BI-1^1/1^ mice (Fig. 4B, Supplementary Table 2).

BI-1 regulates cytosolic Ca^{2+}, leading to the regulation of MAO-A activity. It is known that ER Ca^{2+} release induced by BI-1 can cause reduced intra-ER Ca^{2+} concentrations^{2,5}. We generated HT22 hippocampus neuron cells stably expressing BI-1 or a neomycin-resistant vector control (Neo). The stable expression of BI-1 was confirmed by measuring BI-1 production from BI-1 transfected clones #2-, #8- and #12 (Fig. 5A, upper). Neo-resistant (#4) and BI-1-expressing (#2-, #8- and #12) cells were treated with 1 μM thapsigargin to examine the effects of increased Ca^{2+} availability on MAO-A activity in living cells. Using the cells stably expressing BI-1, we were able to investigate the regulatory effects of BI-1 on cytosolic Ca^{2+} concentration. Although a low level of expression of BI-1 (clone #2) slightly inhibited the Ca^{2+} spike from thapsigargin compared with Neo-resistant vector-transfected (Neo) cells, higher expression of BI-1 (clone #12) significantly inhibited thapsigargin-induced Ca^{2+} stimulation compared with Neo cells. In Figure 5B, higher concentrations of thapsigargin (2 and 5 μM) were used to treat the cells. Compared with the Ca^{2+} increase in Neo cells, the Ca^{2+} increase was significantly regulated in the BI-1 cells compared with Neo cells, and was more significantly regulated in the cells with the highest expressions of BI-1 (#12). The relationship between MAO-A activity and intracellular Ca^{2+} was then analyzed. The thapsigargin-treated BI-1 cells exhibited low levels of MAO-A activity relative to the levels observed in Neo cells (Fig. 5C). BI-1#12 cells were chosen to explore MAO-A activity because of their high level of BI-1 expression. ROS accumulation was examined through DCF-DA staining analyses. A smaller amount of ROS accumulated in thapsigargin-treated BI-1#12 cells compared with control cells (Fig. 5D). Expression of the MAO-A gene (expressed as a ratio to GAPDH) was not affected by thapsigargin treatment (Supplementary Fig. 4A). MAO-A protein expression was also unchanged compared with control levels (Supplementary Fig. 4B), indicating that Ca^{2+} affects MAO-A activity, but not its transcription or translation.

Discussion
The findings of this study demonstrated the endogenous role of BI-1 in resilience against stress, and suggested that BI-1 exhibits...
Ca^{2+}-associated MAO-A regulatory activity. BI-1−/− mice showed vulnerability to chronic mild stress with alterations in hippocampus size and morphology and enhanced ROS accumulation and ER stress response. MAO-A was highly activated, and the concentrations of monoamines including serotonin, norepinephrine and dopamine were reduced, especially following chronic exposure to mild stress. Through an in vitro study, we showed that BI-1 regulates Ca^{2+} release and MAO-A activity, consistent with the in vivo results.

Based on our study, BI-1 gene deficiency is implicated in depression-like behaviors during chronic mild stress. The BI-1−/− stress animals consumed smaller amounts of sucrose than the BI-1+/+ stress group (Fig. 1B), suggesting that BI-1 gene deficiency is involved in the induction of anhedonic-like behavior by mild stress. Compared with the BI-1+/+ stress group, the BI-1−/− stress group showed a significant increase in spontaneous locomotor activity and immobility time (Fig. 1C), suggesting that BI-1 deficiency causes enhanced vulnerability to chronic mild stress. However, it should be noted that younger mice (13 weeks) showed a decrease in only sugar consumption, but not locomotor activities, in BI-1 knockout mice compared with wild-type mice during 1 or 2 weeks of treatment. There may be an age-dependent vulnerability to chronic stress because the animals used in our model were 7–8 months old, relatively older than those used in the previous study. Consistent with our results, the elderly are more susceptible to stress conditions including psychological stress. Oxidative stress has been implicated in a number of age-related degenerative processes in the central nervous system. Why the incidence of chronic disease, including psychiatric diseases, increases with age is still an unanswered question; however, the role of ER stress has recently been a focus in studies of aging processes. The present study suggests that older age may be related to high susceptibility to chronic stress, especially in the BI-1 knockout condition, most likely due to alterations in ER function and ER stress responses.

The BI-1 protein has recently been established as a regulator of ER stress. The ER is a critical site for intracellular Ca^{2+} storage as well as protein synthesis, folding and trafficking. ER stress results in cell death by causing a disturbance of Ca^{2+} homeostasis and leading to activation of caspase-3. ER stress is involved in neurodegenerative diseases such as Parkinson’s disease, Alzheimer’s disease, cerebral ischemia and depression. Notably, certain antidepressants modulate the ER stress response and Ca^{2+} homeostasis. Many antidepressants such as fluvoxamine, a selective serotonin-reuptake inhibitor and imipramine, a tricyclic antidepressant, regulate ER stress. Without the endogenous ER stress regulator gene BI-1, an
Figure 5 | BI-1 regulates Ca\(^{2+}\), leading to the regulation of MAO-A activity. (A) Immunoblotting was performed with anti-HA or anti-β-actin antibody in Neo and BI-1 transfected cells (clones Neo #4, #15 or #34 and BI-1 #2, #8 or #12). Transfected cells were treated with 1 μM thapsigargin and Ca\(^{2+}\) was analyzed as described in the Materials and Methods (lower). (B) Neo and BI-1 expressing clones #2, #8 or #12 cells (expressing difference levels of BI-1) were treated with 2 or 5 μM thapsigargin and Ca\(^{2+}\) was analyzed and quantified. Ca\(^{2+}\) levels in BI-1 cells were significantly different from those in Neo cells for treatment with 2 μM thapsigargin (*p < 0.05) or 5 μM thapsigargin (*p < 0.05). (C) Neo and BI-1 cells were treated with 1, 2 or 5 μM thapsigargin for 30 min and the MAO activity was measured as described in Materials and Methods. (D) Neo and BI-1 cells were treated with 1, 2 or 5 μM thapsigargin for 12 hours and ROS was measured as described in Materials and Methods. ROS levels in BI-1 cells were significantly different from those in Neo cells for each condition (*p < 0.05).
increase in active caspase-3 was detected under chronic mild stress, indicating induction of the ER stress response (Fig. 3B, C). Similarly, overexpressing BI-1 regulates ER stress through the Ca²⁺ buffering capacity of the ER and has been shown to be neuroprotective in several studies. More recently, a transgenic mouse model with BI-1 expression under the control of the neuron-specific enolase promoter showed that BI-1 plays a preventive role against chemical serotonin deprivation stress, although this model did not show any positive ER stress phenomena. However, the results of the study suggested the ability of BI-1 to maintain Ca²⁺ homeostasis and contribute to the preventive/regulatory effect against serotonin deprivation. Homeostatic control of ER Ca²⁺ is essential for appropriate ER function, including protein folding. Elevated cytosolic Ca²⁺ levels have been associated with apoptosis and necrosis in neuronal cultures and this Ca²⁺ disruption may underlie pathologic mechanisms of psychiatric disorders. Considering the pathologic roles of Ca²⁺ disturbance, BI-1-associated Ca²⁺ maintenance may contribute to controlling ER stress-associated progression of depression.

In the BI-1 knockout condition, MAO-A activity induced by chronic stress was significantly higher than that of BI-1 wild-type groups (Fig. 4A), which was correlated with monoamine concentrations (Fig. 4B). In Neuro cells, Ca²⁺ levels increased after treatment with thapsigargin, a Ca²⁺-ATPase inhibitor. The Ca²⁺ increase was smaller in the thapsigargin-treated BI-1 cells compared with the Neuro cells (Fig. 5A). This finding also correlated with MAO-A activity in the thapsigargin-treated condition (Fig. 5C). The requirement of Ca²⁺ for MAO-A activity has recently been demonstrated ex vivo in the brain and is supported by the ability of the Ca²⁺-channel antagonist nimodipine to block the selective increase in MAO-A activity observed in senescence-accelerated mouse brains.

Consistent with this, the Ca²⁺-dependent processes and pathology of encephalopathic brains are linked to a selective increase in MAO-A activity. It has also been suggested that increased intracellular free Ca²⁺ levels and associated MAO-A activity contribute to the oxidative stress associated with neuron degenerative diseases. MAO-A is a risk factor in neuron degenerative diseases such as Alzheimer’s disease. In BI-1 knockout mice, ROS accumulation during catecholamine metabolism due to chronic mild stress (Fig. 2D). Consistent with this, ROS accumulation was much lower in the thapsigargin-treated BI-1-overexpressing neuronal cells compared with control cells (Fig. 5D). Given that 5-HT is metabolized by MAOs and H₂O₂ is generated in the process, the lack of BI-1-associated enhancement of MAO-A activity may increase 5-HT metabolism, resulting in the accumulation of H₂O₂. The mitochondria-bound enzyme MAO generates peroxynitric acids as a natural by-product of the deamination of neurotransmitters such as serotonin. Changes in MAO-A activity parallel changes in the production of ROS such as H₂O₂. Our results also suggest that the oxidative stress associated with chronic mild stress could be due to Ca²⁺-sensitive MAO-A-mediated phenomena, which are regulated by the presence of BI-1.

In conclusion, our findings provide mechanistic support for the hypothesis that BI-1 targets ER Ca²⁺ homeostasis to regulate monoamines including serotonin, norepinephrine and dopamine. These findings hold exciting promise for the development of new treatments for mood disorders. New treatments that specifically target neuronal BI-1 and its regulated pathways may provide innovative mechanisms to enhance brain plasticity in patients with psychiatric disease.

Methods

Reagents and antibodies. 5-Hydroxytryptamine (5-HT), β-phenylethylamine (PEA), the β-actin antibody and protease inhibitor cocktail were bought from Sigma-Aldrich Co. (pH 5-HT (NEC-225) and [14C] PEA (NEC-502) were purchased from PerkinElmer Life Sciences. Phosphate-buffered saline (PBS) was purchased from Invitrogen (Carlsbad, CA). Serotonin, dopamine and norepinephrine were obtained from Sigma (St Louis, MO). Water was purified by a Milli-Q system from Millipore (Bedford, MA). LC-MS grade acetoneitrile, and formic acid were from Fischer Scientific (Fair Lawn, NJ). All other chemicals were of analytical grade and were purchased from Sigma.

Animals. BI-1−/− and BI-1+/− C57BL female mice (n = 33 each, 7–8 months old) were kindly provided by Dr. John C. Reed of the Stanford-Burnham Institute for Medical Research (La Jolla, CA, USA) and housed in groups of 3 to 5 animals at 21 ± 2°C in a 12-hour light/12-hour dark cycle. Mice were maintained in specific pathogen-free housing and were cared for in accordance with the National Institutes of Health Guidelines for Animal Care. The Animal Care and Use Committee of Chonbuk National University Graduate School of Medicine approved this study (Approval number: CBU 2013-0013).

Study design for the chronic mild stress procedure. The procedure for inducing chronic mild stress (CMS) was based on a protocol described previously, with but a difference in mouse age. Briefly, the CMS procedure was applied to each mouse and consisted of a variety of unpredictable mild environmental, social and physical stressors, including confinement in a small tube (1 hour), an empty cage without food (15 hours), water and food deprivation (15 hours), food restriction (approximately 50 mg of food pellets, 3 hours), cage tilted at 45° (1–3 hours), damp sawdust (approximately 200 ml of water per 100 g of sawdust, 3 hours), paired housing in damp sawdust (18 hours), reversal of the light/dark cycle, 4-hour light/ dark succession every 30 minutes (09:00–11:30), light/dark succession every 15–17 hours (9 pm–3 am), 4–6 hours). Control animals were single-housed during the time of CMS to match the condition, but otherwise were group-housed in a different room and left undisturbed with the exception of general handling (e.g., regular cage cleaning and measuring body weight). Female mice aged 7 to 8 months were assigned to eight different groups: BI-1−/− control and stress groups (n = 8–9) and BI-1−/+ control and stress groups (n = 8–9) for either a 2- or 6-weeks period.

Sucrose consumption test. Sucrose intake was measured in the control and stress groups once per weeks on separate days during a 15-hours window (18:00–09:00) following 8 hours of food and water deprivation as previously described.

Consumption was measured by weighing the bottle at the beginning and end of the test. Intake was expressed in relation to animal body weight (g/g).

Spontaneous locomotor activity. Locomotor activity was measured in an open field (40 cm × 50 × 30 cm) consisting of a white base and black walls using a Smart video tracking system (Panlab S.L., Barcelona, Spain) in a softly illuminated (40 W) soundproof room. Each mouse was placed in the center of the open field apparatus and the distance traveled (cm), locomotor time (seconds), number of times crossing the center (50% of all area) and time spent in the center were recorded at 10-min intervals during a 30-min period. The open field arena was thoroughly cleaned between tests.

Magnetic resonance imaging measurements. A total of 124 sagittal T1-weighted (TR = 24, TE = 5, FA = 45) gradient echo images were obtained using a 1.5 T Sigma imager. The MRI volumetric method used was similar to that previously reported.

Sagittal images were acquired with a 24-cm field of view using a slice thickness of 0.1 mm with gaps and a 256 × 192 matrix. For anatomical images, sagittal images were reformatted at a 2-mm slice thickness in the coronal plane. Using the Multimodal Image Data Analysis System (MIDAS) image analysis software (Tsui, M.H. MIDAS Version 1.0, unpublished manual, 1995) developed in-house at NYU with a Sun Sparc workstation (Sun Microsystems, Mountain View, California, hippocampus regions were manually outlined in three-fold enlargement images. All of the image analyses were performed blind to group membership. Tissue volumes were estimated by counting the number of remaining parenchymal pixels (of known size) over the slices measured. The hippocampus and head size were sampled every 2 mm; in most cases this included 12 to 15 coronal sections.

Dihydroethidium staining. Frozen and enzymatically intact sections (30-mm thick) were prepared from mouse brains and immediately incubated with dihydroethidium (10 mol/L; Probes Inc., Eugene, OR, USA) in phosphate-buffered saline for 30 min at 37°C in a humidified chamber. Dihydroethidium is oxidized on reaction with superoxide to ethidium, which binds to DNA in the nucleus to form a fluorescent complex. Images were observed using a fluorescence microscope (PROVISA®X80; Olympus). The intensity of the fluorescence was analyzed and quantified using Image J.

Monoamine oxidase activity. MAO-A and B activity (nmol/hours/mg protein) in BI-1−/− or BI-1−/+ mice and Neomycin resistant vector or BI-1 stably transfect HT-22 cells were measured radiochemically, as previously described. Briefly, for the MAO-A and MAO-B assay, the frozen control or stressed BI-1−/− and BI-1−/+ mouse brain samples were diluted 1 : 3 (vol/vol) with 50 mM potassium phosphate buffer pH 7.5, Aliquots of 100 μL mouse brain homogenate and 250 μL potassium phosphate buffer pH 7.5 were precocubated in a 37°C water bath for 1 min. The enzyme reaction was started by the addition of 150 μL of [14C]-hydroxytryptamine solution (0.03 μCi) for MAO-A and 50 μL of [14C]phenylethylamine solution (0.01 μCi) for MAO-B. After incubating the samples for exactly 15 min at 37°C, the reaction was terminated by addition of 100 μL 4 M HCl. The [14C]-labeled deaminated reaction products 5-hydroxyindolacetic acid and phenylacetic acid were extracted with 5 mL toluene/ethylacetate (1 : 1) by shaking for 10 min on a horizontal shaker (Edmund Bühler SM25, Tübingen, Germany). After centrifugation at 100 × g for 5 min, the aqueous bottom phase was frozen at −70°C for 15 min and the organic phase was decanted into counting vials. After the addition of 5 mL scintillation liquid, the radioactivity
was measured by a liquid scintillation counter. The results were calculated as pmol deaminated metabolite, 5-hydroxyindoleacetic acid and phenylacetic acid formed per min. In both assays, protein content was measured using bovine serum albumin as a standard. For transfected HT-22 cells, the cells were incubated with 1, 2 or 5 μM thapsigargin for 30 min, and then washed with cold PBS and homogenized with 50 mM potassium phosphate buffer pH 7.5. An aliquot (100 μg/100 μL) was incubated with 150 μL of [14C]-hydroxytryptamine (0.03 μCi) for MAO-A and 50 μL [3H]phenylethylamine solution (0.01 μCi) for MAO-B for 10 min at 37°C. All the other processes were the same as for the MAO activity analysis for the brain tissues described above.

Monoamine analysis. Quantification of dopamine, norepinephrine, and serotonin in brain samples was conducted by HPLC coupled with mass spectrometry. HPLC separation was performed using an Agilent 1100 system (Agilent, Palo Alto, CA, USA). Chromatography separation was performed using a ZORBAX SB-Phenyl Narrow Bore column (150 x 2.1 μm internal diameter and 5-μm particle size). Dopamine, norepinephrine and serotonin were separated by gradient elution. The mobile phase was composed of solvent A (water) and solvent B (acetonitrile) and both solvents contained 0.1% formic acid. The gradient run started with 5% solvent B for 1 min and increased to 100% over 20 min, where it remained for another minute before returning to 5% over 25 min, which was maintained for 7 min. The total run time was 32 min at a flow rate of 0.2 mL/min. An Agilent Technologies 6410 triple quadrupole mass spectrometer equipped with electrospray ionization (ESI) in the positive ionization mode was used. The ESI-MS/MS parameters were set as follows: gas temperature 300°C, gas flow 10 L/min and capillary voltage 4 kV. Samples were analyzed in the multiple reaction monitoring (MRM) and the MRM transition used were m/z 177→160 for serotonin, m/z 170→152.1 for norepinephrine and m/z 154.2→137.1 for dopamine. The fragmentation voltages used were 30 V for dopamine and 50 V for serotonin and norepinephrine. The collision energy was set at 3 eV for norepinephrine and 5 eV for serotonin and dopamine. Data acquisition was performed with MassHunter software.

Cell culture. The immortalized mouse hippocampal HT-22 cell line was obtained from the American Type Cell Collection (ATCC). The cells were stably transfected with the Neomycin-resistant pcDNA3 (Neo) or pcDNA3-BI-1 HA vector (BI-1) using the SuperFect transfection reagent (Qiagen, Crawley, UK). The cells were cultured for 5 weeks in 1 mg/mL G418 (Invitrogen, Carlsbad, CA, USA). Cells were maintained in DMEM/low glucose medium containing 10% fetal bovine serum, 100 IU/mL penicillin G sodium salt and 0.03% glutamine at 37°C and 5% CO2.

Calcium analysis. The procedures for Ca2+ measurement were modified from Kim et al.1 Briefly, the low-affinity fluorescent Ca2+ dye Fura-2/AM (1-(2-(5-carboxyoxazol-2-yl)-6-amino fluorescein)-5-methylaminofluorescein) was used to measure intracellular free Ca2+ (Ca2+i). Changes in [Ca2+i], determined as a ratio of 340/380-nm excitation (512-nm emission) using an integrated spectrofluorometer (Photon Technology International, Birmingham, NJ, USA). Ca2+ concentrations were calculated using the equation [Ca2+i] = Kd(Famax - Fam)/Fs(Fmax - Fs) (Rmax - R), where Kd is a constant value of 229 nM was assumed for the binding of calcium to Fura-2/AM. Rmax and Rfmax were determined in each experiment by the consecutive addition of 30 μM Trition X-100 (Rmax) and 50 mM EDTA (Rfmax). Separately, to determine the relative Ca2+ increase, cytosolic Ca2+ was compared between Neos cells and BI-1 cells as a percentage based upon the area of the cytosolic Ca2+ increase curve for Neos cells treated with 2 μM thapsigargin (considered as 100%).

Dichlorofluorescein diacetate (DCF-DA) analysis. The intracellular ROS levels were measured as described previously29. Briefly, 1.5 x 105 cells were incubated with 1, 2 or 5 μM thapsigargin for 12 hours and then treated with 1 μM DCF-DA at 37°C for 30 min. After cooling to ice, the cells were washed with cold phosphate buffered saline (PBS), scraped from the plate and resuspended in PBS containing 10 mM EDTA. The fluorescence intensity of the 2,7'-dichlorofluorescein formed by a reaction between DCF-DA and the intracellular ROS was analyzed for more than 10,000 viable cells from each sample by FACS flow cytometry (Partec, Münster, Germany). An excitation of 488 and emission wavelengths of 525 and 580 nm (FlowJo). The data were collected and analyzed using FlowJo software. All experiments were repeated at least eight times, with similar results. The data were expressed as representative histograms from eight independent experiments.

Statistical analysis. All data are expressed as the mean ± standard error of the mean. Body weight and sucrose consumption were analyzed with a repeated-measures two-way analysis of variance (ANOVA). Individual measures of body weight, sucrose consumption and locomotor activity were compared among the groups using a one-way ANOVA followed by Tukey's post hoc test. Differences of p < 0.05 were considered significant.

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Acknowledgments

This study was supported by National Research Foundation (2012R1A2A1A001907 and 2008-006279) and by Korea Healthcare Technology R&D Project (A121931), Ministry for Health and Welfare, Republic of Korea. We thank Professor Myoung-Goo Kang, at Korea Advanced Institute of Science and Technology, for critical review of this manuscript.

Author contributions

H.Y.L., G.H.L. and T.R.B. performed experiments and analyzed data; M.A. analyzed monoamines, S.M.L. measured brain micro-CT image, M.R.L. analyzed data, K.Y.J. and H.H.L. reviewed the manuscript and H.R.K. and H.J.C. designed the research, interpreted data and edited the manuscript. All authors reviewed the manuscript.

Additional information

Supplementary information accompanies this paper at http://www.nature.com/scientificreports

Competing financial interests: The authors declare no competing financial interests.

How to cite this article: Lee, H.-Y. et al. The protective role of Bax Inhibitor-1 against chronic mild stress through the inhibition of monoamine oxidase A. *Sci. Rep.* 3, 3398; DOI:10.1038/srep03398 (2013).

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