

## NIH Public Access

Author Manuscript

*Neurobiol Aging*. Author manuscript; available in PMC 2013 June 1

Published in final edited form as:

Neurobiol Aging. 2012 June ; 33(6): 1125.e9–1125.e18. doi:10.1016/j.neurobiolaging.2011.11.023.

# Calpastatin modulates APP processing in the brains of $\beta$ -amyloid depositing but not wild-type mice

Jose Morales-Corraliza<sup>1,2</sup>, Jason D Berger<sup>1</sup>, Matthew J Mazzella<sup>1</sup>, Veeranna<sup>1,2</sup>, Thomas A. Neubert<sup>2</sup>, Jorge Ghiso<sup>2</sup>, Mala V Rao<sup>1,2</sup>, Matthias Staufenbiel<sup>3</sup>, Ralph A Nixon<sup>1,2</sup>, and Paul M Mathews<sup>1,2</sup>

<sup>1</sup>Nathan Kline Institute for Psychiatric Research, Orangeburg, NY 10962, USA <sup>2</sup>New York University School of Medicine, New York, NY 10016, USA <sup>3</sup>Novartis Institutes for Biomedical Research, Nervous Systems Research, Basel CH-4056, Switzerland

#### Abstract

We report that neuronal overexpression of the endogenous inhibitor of calpains, calpastatin (CAST), in a mouse model of human Alzheimer's disease (AD) β-amyloidosis, the APP23 mouse, reduces  $\beta$ -amyloid pathology and A $\beta$  levels when comparing aged, double transgenic (tg) APP23/ CAST with APP23 mice. Concurrent with A $\beta$  plaque deposition, aged APP23/CAST mice show a decrease in the steady-state brain levels of the amyloid precursor protein (APP) and APP Cterminal fragments when compared to APP23 mice. This CAST-dependent decrease in APP metabolite levels was not observed in single tg CAST mice expressing endogenous APP or in younger, AB plaque predepositing APP23/CAST mice. We also determined that the CASTmediated inhibition of calpain activity in the brain is greater in the CAST mice with  $\beta$ -amyloid pathology than in non-APP tg mice, as demonstrated by a decrease in calpain-mediated cytoskeleton protein cleavage. Moreover, aged APP23/CAST mice have reduced ERK1/2 activity and tau phosphorylation when compared to APP23 mice. In summary, in vivo calpain inhibition mediated by CAST transgene expression reduces A $\beta$  pathology in APP23 mice, with our findings further suggesting that APP metabolism is modified by CAST overexpression as the mice develop  $\beta$ -amyloid pathology. Our results indicate that the calpain system in neurons is more responsive to CAST inhibition under conditions of  $\beta$ -amyloid pathology, suggesting that in the disease state neurons may be more sensitive to the therapeutic use of calpain inhibitors.

#### Keywords

calpain; calpastatin; APP; Aβ; Alzheimer's disease

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Correspondence to be addressed to: J. Morales-Corraliza and P.M. Mathews, Nathan Kline Institute and New York University School of Medicine, Orangeburg, New York, 10962 (USA), Tel 845 398 5439, Fax 845 398 2197, jmorales-corraliza@nki.rfmh.org, mathews@nki.rfmh.org.

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#### Introduction

Disruption of multiple proteolytic systems contributes to Alzheimer's disease (AD) pathobiology, including alterations in the enzymes responsible for  $\beta$ -amyloid (A $\beta$ ) generation and clearance and dysfunction of the lysosomal system (Mathews, et al., 2002a, Nixon, et al., 2000). Calpains have also shown evidence of hyperactivity in human AD tissue (Grynspan, et al., 1997, Liu, et al., 2005, Nixon, 2003, Saito, et al., 1993). The calpain family is a group of Ca<sup>2+</sup>-activated, cytosolic, neutral pH, cysteine proteases (Huang and Wang, 2001) which modulate, probably indirectly, the localization and proteolytic processing of the amyloid precursor protein (APP) in cultured cells (Mathews, et al., 2002b). The most abundantly expressed calpains are m-calpain and µ-calpain, which are distinguished by their different affinities for  $Ca^{2+}$ , each of which forms a functional heterodimer with the shared regulatory calpain small subunit 1 (Capn4) (Croall and DeMartino, 1991, Sorimachi, et al., 1997). In addition to their Ca<sup>2+</sup>-dependence, calpains are regulated by cytosol-to-membrane translocation and an endogenous inhibitor, calpastatin (CAST). The CAST protein consists of four calpain-inhibitory domains that are subjected to calpain cleavage and terminal inactivation by caspase (Croall and DeMartino, 1991, Goll, et al., 2003, Maki, et al., 1991, Sorimachi, et al., 1997, Wang, et al., 1998).

Activation of calpains in AD is evidenced by increased levels of activated  $\mu$ -calpain (Saito, et al., 1993) and m-calpain (Grynspan, et al., 1997) in neurons. The role of the calpain system in normal brain function and in pathological conditions has also been examined in various mouse models with modified calpain and CAST expression. While genetic deletion of either the m-calpain large subunit or the single calpain small subunit is lethal (Arthur, et al., 2000, Takano, et al., 2005), deletion of the  $\mu$ -calpain large subunit does not result in an apparent gross phenotype (Azam, et al., 2001, Grammer, et al., 2005). Mice lacking CAST, while showing decreased locomotor activity and a decreased acoustic startle response, have no change in hippocampal-dependent memory function (Nakajima, et al., 2008). CAST overexpressing mice similarly do not have gross memory deficits (Higuchi, et al., 2005, Takano, et al., 2005), arguing that CAST deletion or overexpression produces, in general, mild effects in the normal mouse brain (Rao, et al., 2008).

The role of the calpain system in AD pathobiology has been recently explored in  $\beta$ -amyloid depositing mouse models. In Tg2576 (Vaisid, et al., 2007) and in APP/PS1 mice (Liang, et al., 2010), calpain appears to be activated and CAST diminished, consistent with reports in human AD tissue (Grynspan, et al., 1997, Liu, et al., 2005, Nixon, 2003, Saito, et al., 1993). In a mouse overexpressing wild-type APP that does not develop  $\beta$ -amyloid pathology, calpains also appear to be activated in neurons (Kuwako, et al., 2002). In APP/PS1 mice, chronic calpain inhibition has been shown to reduce amyloid plaque burden and improve memory and synaptic transmission (Liang, et al., 2010, Trinchese, et al., 2008). This is in contrast to previous results from multiple laboratories showing that the acute pharmacological inhibition of calpains in cell culture systems dramatically increases A $\beta$ 42 generation (Klafki, et al., 1996, Mathews, et al., 2002b, Yamazaki, et al., 1997, Zhang, et al., 1999), which is thought to be a more pathological A $\beta$  species. Here, we examined brain APP metabolite levels in mice overexpressing CAST in neurons compared to wild-type mice, as well as CAST overexpressing mice crossed to the β-amyloid depositing APP23 line. While CAST-overexpression-mediated changes were not seen in otherwise wild-type mice, the development of  $\beta$ -amyloid pathology in the APP23 mice corresponded to both an increased sensitivity of the calpain system to CAST inhibition and a reduction in APP metabolite levels.

#### Materials and methods

#### Transgenic mice

All experiments involving mice received prior approval from the Nathan Kline Institute Animal Care and use committee. Neuron-specific overexpression of human CAST is driven by the Thy-1.2 promoter as previously described (Rao, et al., 2008). APP23 mice, which overexpress Swedish-mutant human APP (K670N, M671L) under the Thy-1.2 promoter, develop  $\beta$ -amyloid plaque pathology after the first year of life (Sturchler-Pierrat, et al., 1997). The depositing APP23/CAST and APP23 mice used in this study were female 13month old mice. All mice were maintained on a C57B16 background.

#### Brain Processing, Western blotting and Aß ELISA measurements

Ten-percent (weight-to-volume) homogenates were prepared from a hemibrain lacking the olfactory bulb and cerebellum, and used for biochemical analyses (Schmidt, et al., 2005a). An aliquot of the homogenates was extracted in diethylamine (DEA) prior to sandwich ELISA to detect soluble murine  $A\beta$  in single tg CAST and wild-type mice as well as soluble human  $A\beta$  in the predepositing APP23/CAST and APP23 mice (Schmidt, et al., 2005b). DEA-extracts of all genotypes were also used for sAPP isolation prior to Western blot analyses as previously described (Morales-Corraliza, et al., 2009). In aged APP23/CAST and APP23 mice, an aliquot of the homogenate was extracted in formic acid for  $\beta$ -amyloid-associated  $A\beta$  quantitation (Schmidt, et al., 2005a, Schmidt, et al., 2005b). For Western blotting, proteins were sized by SDS-PAGE, transferred to polyvinylidene difluoride membrane (Mathews, et al., 2002b), and incubated with antibodies as previously described (Morales-Corraliza, et al., 2009). For unbiased analysis of  $A\beta$  species,  $A\beta$  in brain homogenates was resolved by urea/SDS-PAGE as previously described (Klafki, et al., 1996) prior to Western blotting analysis.

#### Antibodies

Antibody C1/6.1 recognizes the carboxyl-terminal cytoplasmic domain of APP (Mathews, et al., 2002b), and m3.2 recognizes residues 10–15 of murine Aβ, also detecting murine APP, sAPPα and Aβ (Morales-Corraliza, et al., 2009). 22C11 was purchased from Millipore (Temecula, CA) and detects APP N-terminal in both human and mouse epitopes. Monoclonal antibody 6E10 (Covance, Princeton, NJ), which recognizes residues 1–16 of human A $\beta$ , was used to detect human sAPP $\alpha$  and A $\beta$ . The antibody 4G8 (Covance, Princeton, NJ) recognizes residues 17–24 of A $\beta$  and was used to detect A $\beta$  and APP. Mouse monoclonal antibody 6A1 detects sAPPß containing the Swedish mutation but not the wildtype sequence (Morales-Corraliza, et al., 2009). Neprilysin was detected with the monoclonal antibody 56C6 (CD10) (Novacastra, Newcastle, UK), and IDE with the rabbit polyclonal antibody IDE1 ((Qiu, et al., 1998); a gift of Dr. Dennis Selkoe). Cytoskeletal proteins and kinases were detected with antibodies against MAP1 (Clone HM-1; Chemicon, Temecula, CA), MAP2 (antibody 18-1; (Rao, et al., 2008)) and all-spectrin (MAB1622; Chemicon, Temecula, CA), CDK5 (C-8:sc-173; Santa Cruz Biotechnology, Santa Cruz, CA), GSK3β (27C10; Cell Signaling Technology, Danver, MA), and ERK1/2 and Phospho-ERK1/2 (9102 and 9101; Cell Signaling Technology, Danver, MA). Phosphorylated tau was detected using PHF1 (Chemicon International, Temecula, CA; phospho-epitope at Ser396/404) and CP13 (a gift of Dr. Dennis Selkoe; phospho-epitope at Ser202), while T57120 (BD Biosciences, San Jose, CA), MN37 (a gift of Dr. Dennis Selkoe) and tau1 (Chemicon International, Temecula, CA) were used to detect total tau independent of its phosphorylation state. Phosphorylated  $\alpha$ -synuclein was detected using a phospho-Ser129specific a-synuclein antibody (Abcam, Cambridge, MA). Syn-1 antibody (BD Biosciences, San Jose, CA) was used for phospho-independent  $\alpha$ -synuclein recognition. Activated calpain II immunolabeling was done using the rabbit polyclonal antibody C24 (Rao, et al., 2008).

#### Thioflavin S Staining

 $\beta$ -amyloid plaque burden was visualized by florescence microscopy of 35-µm thick vibratome sections prepared from formalin-fixed hemibrains previously stained with Thioflavin S as described in (Mi, et al., 2007). Thioflavin labeling was quantitated by the program AxioVision 4.6 (n=5 in APP23/CAST and n=5 in APP23 mice; 5 sections per mouse).

#### Immunocytochemistry

Vibratome sections were blocked with 10% horse serum (Invitrogen, Carlsbad, CA) at room temperature and incubated with the primary antibody C24 (Rao, et al., 2008) overnight at 4° C. Sections were immunostained using a biotinylated goat anti-rabbit secondary antibody and Vectastain ABC kit (both products, Vector Laboratories, Burlingame, CA).

#### Matrix-Assisted Laser Desorption/Ionization Time-of-Flight (MALDI-TOF) Mass Spectrometry

Brain homogenates were immunoprecipitated with a combination of 6E10 and 4G8 antibodies (Covance, Princeton, NJ) using Dynabead M-280 sheep anti-mouse IgG (Invitrogen, Carlsbad, CA) (Tomidokoro, et al., 2010). Proteins were eluted with C4 ZipTip (Millipore, Billerica, MA) in 90% acetonitrile and 0.1% trifluoracetic acid (TFA), mixed 50:50 (v:v) with 10 mg/ml  $\alpha$ -Cyano-4-hydroxycinnamic acid in 0.1% TFA and 50% acetonitrile, and air dried. Samples were analyzed using a Bruker Autoflex MALDI-TOF mass spectrometer (Bruker Daltonics, Billerica MA) in positive ion linear mode using standard operating conditions at the NYU Protein Mass Spectrometry Core for Neuroscience.

#### **Statistical Analysis**

Western blots were quantitated using ImageJ (http://rsb.info.nih.gov). ELISA measurements were analyzed using a non-parametric Mann-Whitney u test. All data were plotted on GraphPad Prism v.5 (GraphPad Software, San Diego, CA, USA) for statistical analysis. Throughout, results are expressed as the mean ± SEM.

#### Results

To explore the relationship between neuronal calpain function and AD pathology, we crossed human CAST overexpressing tg mice (showing ~7-times the levels of the endogenous CAST expression in the brain (Rao, et al., 2008)) with an amyloid precursor protein (APP) overexpressing tg line that develops  $\beta$ -amyloid plaques (APP23 mice; (Sturchler-Pierrat, et al., 1997)). Brain plaque pathology in littermate 13-month-old female APP23/CAST and APP23 double tg was visualized by Thioflavin S staining, with the area occupied by plaque in the double tg mice decreased by ~60% (plaque area as % of total brain: 0.13%±0.02 in APP23/CAST mice versus 0.34%±0.03 in APP23 mice; p<0.001) (Figure 1A-B). ELISA measurements of formic acid-extractable AB in the APP23/CAST double tg versus the APP23 single tg mice showed a decrease of 70%±11 for deposited human A $\beta$ 42 (p<0.01) and a similar decrease for human A $\beta$ 40 levels (77%±15; (p<0.01)) (Figure 1C). Similar decrease in co-deposited murine A $\beta$ 40 and A $\beta$ 42 was also observed by sandwich ELISA (decrease of  $61\% \pm 22$  in A $\beta$ 42 (p<0.05) and  $63\% \pm 22$  in A $\beta$ 40 (p<0.05); Supplemental Figure 1A). Thus, the A $\beta$  burden is reduced in aged APP23 mice overexpressing CAST, with the ratio of Aβ42 to Aβ40 remaining similar (APP23/CAST: 4.7±0.3 and APP23: 4.1±0.5; p>0.05).

Since inhibiting calpain activity has been shown to change the ratio of A $\beta$ 40 and A $\beta$ 42 generated by APP overexpressing cells in culture (Klafki, et al., 1996, Mathews, et al.,

2002b, Yamazaki, et al., 1997, Zhang, et al., 1999), we examined A $\beta$  in these mice using urea SDS-PAGE in order to detect all A $\beta$  species (Figure 1D). In agreement with our  $\beta$ -amyloid ELISA findings (Figure 1C), we observed a decrease in A $\beta$  band densities in APP23/CAST versus APP23 mice, with the relative ratios of the three predominant A $\beta$  species (A $\beta$ 38, A $\beta$ 40 and A $\beta$ 42) not showing changes. Additionally, we analyzed A $\beta$  peptides by MALDI-TOF mass spectrometry. While this technique does not allow for the direct quantification of different A $\beta$  species (Tomidokoro, et al., 2010), the ratio between the peak intensity of the A $\beta$  species reflects the relative abundance of the peptides. No differences in the relative abundance of A $\beta$  species were seen when comparing APP23/CAST with APP23 mice (Figure 1E), consistent with the ELISA measurements (Figure 1C) and the urea SDS-PAGE analysis (Figure 1D).

Given this decrease in  $\beta$ -amyloid in the APP23/CAST mice, we next determined the levels of two A $\beta$  degrading enzymes within the brain: neprilysin and insulin degrading enzyme (IDE). No changes in protein levels in these proteases were observed when comparing APP23/CAST with APP23 mice (Figure 1F). APP levels in the brain of 13-month-old A $\beta$ plaque depositing APP23/CAST mice, however, showed a decrease of 25%±7 compared to APP23 mice ((p<0.05); Figure 1G–H). Consistent with this reduction in APP, APP Cterminal fragments (CTF) levels were also decreased in APP23/CAST mice by 36%±12 ((p<0.05); Figure 1G–H). We determined that the highly stable secreted APP (sAPP) metabolites (Morales-Corraliza, et al., 2009) showed no changes between APP23/CAST and littermate APP23 mice, either when probing with an antibody that detect both sAPP $\alpha$  and sAPP $\beta$  (sAPP total) or antibodies that only detect sAPP $\alpha$  or sAPP $\beta$  (Figure 1G). Relative levels of APP, CTFs and sAPP are shown graphically in the Figure 1H.

We extended our study on CAST overexpression in vivo by examining APP metabolite levels in CAST mice without Aβ-plaque deposition by examining younger (4-month old), predepositing APP23/CAST mice compared to APP23 mice and by comparing CAST single tg to wild-type mice. Predepositing APP23/CAST mice did not show a significant change in human A $\beta$  brain levels compared to littermate APP23 mice (Figure 2A). At this age, no changes in the levels of APP, CTFs, sAPP total, sAPP $\alpha$  and sAPP $\beta$  were seen (Figure 2B), which is in contrast to the reduction in APP and CTF levels in the 13-month-old mice (Figure 1G–H). This suggests that calpastatin expression mediated changes in neuronal APP processing are not prominent until the mice are older and beginning to show brain  $\beta$ -amyloid pathology. To further examine this result, we analyzed at various ages APP metabolite levels in CAST mice expressing only the endogenous murine APP and therefore without  $A\beta$ accumulation or plaque pathology. At multiple ages from 4 to 24 months in single CAST tg mice (shown in Figure 2C-D are findings from 18-24 month old mice; other ages are shown in Supplement Figure 1B–D), no changes were seen in APP metabolite levels – including APP, CTFs, sAPP total, sAPP $\alpha$  and A $\beta$  – when compared to littermate non-tg mice (Figure 1I–H).

By Western blot analysis, we also characterized in APP23/CAST and APP23 mice, calpainmediated cleavage of cytoskeleton proteins, including MAP1 and MAP2, and  $\alpha$ II-spectrin (Fifre, et al., 2006, Pike, et al., 2001, Warren, et al., 2007). In Figure 3A, Western blot analysis of the total protein and calpain-specific breakdown products of MAP1 (Fifre, et al., 2006) showed lesser amounts of the breakdown products in APP23/CAST mice of either age when compared to APP23 littermates. Total MAP2 protein levels remaining in APP23/ CAST were found to be greater than the levels seen in APP23 littermates, and these changes in MAP2 levels were not as great in single tg CAST compared to non-tg mice. Figure 3B shows that while  $\alpha$ II-spectrin cleaved by caspase showed no significant changes between the two groups, calpain-mediated  $\alpha$ II-spectrin breakdown product (Pike, et al., 2001, Warren, et al., 2007) levels were reduced in the APP23/CAST mice compared to single APP23 tg mice.

Figure 3C shows a graphic representation of the analysis of the band density of total proteins and breakdown products of MAP2, MAP1 and  $\alpha$ II-spectrin blots. Significant decreases in calpain activity were found in APP23/CAST and CAST mice compared to APP23 and nontg mice, respectively. We also found greater CAST-mediated calpain inhibition when comparing APP23/CAST to APP23 mice than when CAST single tg were compared to nontg mice (an apparent decrease in calpain activity of 41%±10 in APP23/CAST versus APP23 mice and 24%±7 in CAST versus non-tg mice; p<0.05 between the groups). This is consistent with calpain inhibition itself being greater in  $\beta$ -amyloid depositing mice expressing CAST (APP23/CAST versus APP23 mice) compared to CAST single tg versus non-tg mice. In Figure 3D, the immunolabeling pattern seen with an antibody that detects activated calpain II (C24; (Rao, et al., 2008)) is in agreement with our findings above on calpain activity indicated by cytoskeletal protein Western blotting (Figure 3A–C).

The phosphorylation of tau can be driven by A $\beta$  accumulation in the brain (Gotz, et al., 2001, Lewis, et al., 2001), potentially by kinases whose activity can be modulated by calpains (Amadoro, et al., 2006, Hye, et al., 2005, Lee, et al., 2000, Veeranna, et al., 2004). Given the potential for aberrant activity of these kinases in AD, we determined the levels and activity/phosphorylation state of key tau kinases as well as tau using phospho-dependent and phospho-independent antibodies (Figure 4). While the level of the catalytic domain of CDK5 and level of GSK3β did not change, a decrease in phosphorylated ERK1/2 levels (active form compared to non-phosphorylated ERK1/2) was seen in APP23/CAST compared to APP23 mice (Figure 4). Additionally, the binding of two antibodies - PHF1 and CP13 – that recognize tau phospho-epitopes of ERK1/2 was found to be significantly reduced in APP23/CAST animals, consistent with a decrease in phosphorylated ERK1/2 levels. The phospho-independent tau antibodies T57120, MN37 and tau1 showed no differences in total tau levels between APP23/CAST and APP23 mice. This is in contrast to the calpain-mediated turnover of other cytoskeletal proteins as shown in Figure 3A–D, but consistent with the lack of tau-fragments seen in other mouse models of human  $\beta$ amyloidosis (Liang, et al., 2010, Roberson, et al., 2007) and in agreement with the eventual accumulation of tau seen as paired-helical filament in human AD. To expand the potential disease relevance of these studies, we additionally examined the levels of phosphorylated asynuclein (phospho-Ser129  $\alpha$ -synuclein) and total  $\alpha$ -synuclein. No changes in  $\alpha$ -synuclein signal were detected when comparing APP23/CAST with APP23 mice, consistent with the idea that the calpain system is particularly responsive to  $A\beta/\beta$ -amyloid accumulation, and that this appears to have downstream effects on tau phosphorylation.

#### Discussion

The pathological role of calpain hyperactivation following the unregulated increase in intracellular Ca<sup>2+</sup> concentration that accompanies excitotoxicity has long been appreciated (Bartus, et al., 1995, Wang, 2000). This activation of calpains results in the cleavage of a number of neuronal substrates that can negatively affect neuronal structure, function, and survival (Siman and Noszek, 1988). However, the involvement of calpains in mediating neuronal vulnerability in chronic, long-term disease is less well understood. Multiple studies suggest that in post-mortem human AD brain the neuronal calpain system is upregulated (Grynspan, et al., 1997, Liu, et al., 2005, Nixon, 2003, Saito, et al., 1993), albeit these findings do not necessarily argue for a direct role for calpain activation in the disease. There are numerous studies, however, indicating that calpain activation may lead to pathological changes in tau phosphorylation both *in vitro* (Amadoro, et al., 2006, Chung, 2009, Lee, et al., 2000) and *in vivo* (Chung, 2009, Veeranna, et al., 2004), as well as alterations in APP metabolism *in vitro* (Klafki, et al., 1996, Mathews, et al., 2002b, Yamazaki, et al., 1997, Zhang, et al., 1999). The growing evidence from transgenic mouse models, including our study, that calpain activation can result from  $\beta$ -amyloid accumulation and/or altered A $\beta$ 

levels argues that the calpain system responds to AD-related pathological changes in the brain (Liang, et al., 2010, Vaisid, et al., 2007), thus positioning calpains as having a role in both driving and responding to the disease.

We initiated this study because of the evidence from multiple groups, including ours (Klafki, et al., 1996, Mathews, et al., 2002b, Yamazaki, et al., 1997, Zhang, et al., 1999), that the acute pharmacological inhibition of calpains in cell culture systems dramatically increases A $\beta$ 42 generation, suggesting that inhibition of calpains in the brain might lead to greater  $\beta$ -amyloid pathology due to increased A $\beta$ 42 levels. *In vivo*, however, we now report the opposite, with CAST-transgene-overexpression-mediated calpain inhibition leading to a reduction in  $\beta$ -amyloid pathology while having no effect on the ratio of the various A $\beta$  peptides detected in the brain. Indeed, we choose to use a model that develops  $\beta$ -amyloid pathology without artificial A $\beta$ 42-drive, such as the mutant PS1 expression in the APP/PS1 $\Delta$ 9 × CAST tg crosses described by Liang *et al.* (Liang, et al., 2010), in order to be able to detect a CAST-overexpression effect on A $\beta$  C-terminal cleavage. In addition to the evidence we present that there are no changes in the C-terminal cleavage-site of the A $\beta$  derived from the human Swedish APP, we did not detect changes in the murine A $\beta$ 40:A $\beta$ 42 ratio from the endogenous, wild-type APP in either single CAST tg mice or in the APP23/CAST and APP23 mice (see Figure 1 and 2 and Supplemental Figure 1).

We found that in AB depositing APP23/CAST mice, CAST overexpression leads to a decrease in multiple APP metabolite levels, including APP, CTFs, and AB. This reduction in brain APP metabolite levels with CAST overexpression did not occur either in wild-type mice throughout their life-span or in predepositing APP23 mice. Indeed, our findings suggest that only with  $\beta$ -amyloid pathology in the aged APP23 mice, CAST overexpression produces changes in APP metabolism. Additionally, changes in the activity of the calpain system by CAST observed in vivo in this study are consistent with those seen in human AD tissue (Grynspan, et al., 1997, Liu, et al., 2005, Nixon, 2003, Saito, et al., 1993) and in mouse models (Liang, et al., 2010, Vaisid, et al., 2007) subsequent to the formation of  $\beta$ amyloid plaques, where calpains appear to be activated and CAST levels decreased. The influence of APP metabolism on calpain activity was also observed in a previous report in mice overexpressing wild-type APP that showed that, although these mice do not develop  $\beta$ amyloid pathology, calpains are activated in neurons (Kuwako, et al., 2002). In this study, APP-induced calpain activation, which was sensitive to calpain inhibitors in vivo, was not seen in mice expressing an APP mutant that is not processed to produce A<sup>β</sup>. This provides further evidence that calpain activation in these models is dependent upon altered brain  $A\beta$ levels. By examining the levels of calpain substrates and cleavage-products, our findings show that the inhibition of calpain activity by CAST is greater in the aged APP23 mice with  $\beta$ -amyloid pathology (and apparent in the predepositing APP23 mice), when compared to single tg CAST mice, which is consistent with the idea that the calpain system is more responsive to CAST overexpression when perturbed by excess brain AB. That CAST overexpression-mediated modulation of calpain occurs under conditions of A $\beta$  induced neuronal stress is also in agreement with the findings of Rao et al. (Rao, et al., 2008) showing that CAST overexpression inhibits pathologically elevated calpain activity in vivo following excitotoxicity, but less under baseline conditions. Our study supports the idea that, *in vivo*, there is cross-talk between developing Aβ pathology and the calpain/CAST system, which may show elevated specific activity to the disease that may render the calpain system more responsive to inhibition.

Our findings argue that restoring calpain homeostasis has multiple beneficial effects, including reducing  $\beta$ -amyloid accumulation and tau phosphorylation. While neither our study nor prior studies (Chung, 2009, Liang, et al., 2010) differentiate between reductions in tau phosphorylation resulting from less A $\beta$  accumulation or directly from modulation of

calpain activity, either mechanism is potentially beneficial in the disease. Pathologically important disruption of calpain activity, including dysregulation of tissue-specific calpain family members, can occur in a number of aging-related diseases, including type 2 diabetes, cataracts, muscular dystrophy, Parkinson's disease, rheumatoid arthritis, ischemia, stroke and brain trauma, various platelet syndromes, hypertension, liver dysfunction and some types of cancer (Carragher, 2006, Zatz and Starling, 2005). When associated with a specific calpain family member, disease development and progression appears to be directly linked to altered calpain expression and/or activity. Currently, the evidence from mouse models would suggest that changes in calpain activity in AD correlate with developing A $\beta$ pathology. Such vulnerability of the calpain system in AD would appear to offer an opportunity for therapeutic modulation, with the CAST overexpression systems suggesting that inhibition and/or restoration of more normal calpain activity has benefits both by reducing  $\beta$ -amyloid accumulation and potentially by reducing tau phosphorylation.

#### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

#### Acknowledgments

We would like to thank Mr. Steven Blais of the NYU Protein Mass Spectrometry Core for Neuroscience for MALDI-TOF analysis, Drs. Haung Yu and Karen Duff for the kind gift of the synuclein antibodies and Dr. Panaiyur Mohan for advice on the assessment of calpain activity *in vivo*. This work was supported by the Alzheimer's Association (IIRG-07-60047 to P.M.M), the NINDS (NS045205 to P.M.M) and NIA (AG017617 to P.M.M, R.A.N).

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### Figure 1. CAST overexpression decreases APP metabolite levels in depositing APP23/CAST mouse brain. (A–B)

Brain sections of depositing 13-month-old APP23/CAST and APP23 mice were used to visualize and quantify  $\beta$ -amyloid plaque area by Thioflavin S staining. (C) Formic acid-extractable human A $\beta$ 40 and A $\beta$ 42 levels were measured by sandwich ELISA. (D) Analysis of A $\beta$ 38, A $\beta$ 40 and A $\beta$ 42 using Western blots as previously described by Klafki *et al.* (Klafki, et al., 1996) based upon migration of A $\beta$  standards. (E) MALDI-TOF mass spectrometry of brain homogenates immunoprecipitated simultaneously with 6E10 and 4G8 antibodies. (F) Neprilysin and IDE levels are shown by Western blot analysis of brain homogenates. (G) By Western blot, total proteins probed for APP and CTFs; soluble brain extracts lacking membrane-associated proteins probed for sAPP total, sAPP $\alpha$  and sAPP $\beta$ , as indicated (Morales-Corraliza, et al., 2009). (H) Graphic representation of the analysis of the band density of the blots of APP, CTFs and sAPP total. \*p<0.05; \*\* p<0.01; \*\*\*p<0.001.





Soluble human A $\beta$ 40 and A $\beta$ 42 levels and (**B**) additional APP metabolites levels, including APP, CTFs, sAPP total, sAPP $\alpha$  and sAPP $\beta$  in predepositing 4-month-old APP23/CAST and APP23 mouse brain. (**C**) Soluble endogenous A $\beta$ 40 and A $\beta$ 42 levels and (**D**) additional APP metabolite levels in 18-month-old CAST and wild-type mice.



#### Figure 3. Assessment of calpain activity in APP23/CAST and APP23 mouse brain

Characterization of calpain activity in APP23/CAST mice by Western blot analysis of calpain-mediated clearance/cleavage of cytoskeleton proteins (Fifre, et al., 2006, Pike, et al., 2001, Rao, et al., 2008, Warren, et al., 2007) (A) Blots show total protein remaining for MAP1 and MAP2 and the major breakdown products of MAP1. (B) Total protein remaining and the two major breakdown products of  $\alpha$ II-spectrin, including a 120 kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band corresponding to  $\alpha$ II-spectrin cleaved by caspase and 150kDa band c

three blots as well as additional blots. (D) Immunolabeling with activated calpain II-specific antibody (antibody C24) of serial brain coronal sections of APP23 (a), APP23/CAST (b), wild-type (c) and CAST mice (d). \*p<0.05; \*\* p<0.01; \*\*\*p<0.001.



Figure 4. CAST overexpression inhibits ERK1/2 dependent tau phosphorylation in APP23/ CAST mouse brain

Brain proteins levels of tau kinases and tau in 13-month-old depositing APP23/CAST compared to APP23 mice were determined by Western blotting. Levels of the catalytic domain of CDK5 (panel as indicated throughout) and GSK3 $\beta$  and levels of phospho-ERK1/2 (phospho-dependent antibody) compared to total ERK1/2 (probed with a phospho-independent antibody; see Methods). Levels of tau using two phosphorylation-dependent tau antibodies which recognize tau phospho-epitopes for ERK1/2: PHF1 (phospho-epitope at Ser396/404) and CP13 (phospho-epitope at Ser202) and three phospho-independent tau antibodies T57120, MN37 and tau1. Levels of  $\alpha$ -synuclein using a phosphorylation-dependent  $\alpha$ -synuclein antibody (Syn-1).