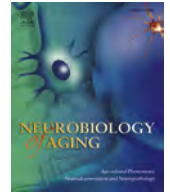




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Age-related memory deficits are associated with changes in protein degradation in brain regions critical for trace fear conditioning



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ABSTRACT

Brain aging is accompanied by an accumulation of damaged proteins, which results from deterioration of cellular quality control mechanisms and decreased protein degradation. The ubiquitin-proteasome system (UPS) is the primary proteolytic mechanism responsible for targeted degradation. Recent work has established a critical role of the UPS in memory and synaptic plasticity, but the role of the UPS in age-related cognitive decline remains poorly understood. Here, we measured markers of UPS function and related them to fear memory in rats. Our results show that age-related memory deficits are associated with reductions in phosphorylation of the Rpt6 proteasome regulatory subunit and corresponding increases in lysine-48 (K48)-linked ubiquitin tagging within the basolateral amygdala. Increases in K48 polyubiquitination were also observed in the medial prefrontal cortex and dorsal hippocampus. These data suggest that protein degradation is a critical component of age-related memory deficits. This extends our understanding of the relationship between the UPS, aging, and memory, which is an important step toward the prevention and treatment of deficits associated with normal cognitive aging and memory-related neurodegenerative diseases.

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1. Introduction

Even in the absence of brain disease, the gradual decline of cognitive ability with age is a growing problem. Age-related cognitive deficits exist on a continuum, wherein some individuals may develop pathologies such as Alzheimer's or Parkinson's disease, and other individuals do not display significant age-related deficits in cognitive ability (Rowe and Kahn, 1987). On the other hand, most individuals age somewhere between these 2 extremes of pathological and successful aging, in a pattern termed normal cognitive aging (Roberson et al., 2012). The proportion of the U.S. population aged over 65 years is projected to more than double by the year 2050, and one in every 5 individuals will be classified as aged (Ortman et al., 2014). Thus, the negative consequences of normal brain aging remain a major challenge to modern neuroscience, while the neurobiology underlying age-related cognitive decline remains largely unknown.

Aging is associated with a variety of biological changes at the cellular level which can be observed in gene expression, intracellular signaling, and metabolism. One critical component of the cellular aging process is a gradual decline in protein homeostasis, which is accompanied by an age-related decrease in function of the ubiquitin-proteasome system (UPS). The UPS is a major regulatory pathway that is responsible for the recognition and clearance of abnormal or damaged proteins, as well as regulation of short-lived proteins in response to intracellular and extracellular signals (Jarome and Helmstetter, 2013). The UPS works through the attachment of polyubiquitin chains to target proteins, and these chains serve as a signal for degradation and proteolysis of the target by proteasomes. Ubiquitin ligases tag proteins for degradation by attaching polyubiquitin chains that identify them for processing by the 26S proteasome. The 26S complex consists of a 20S catalytic core and 2 19S regulatory particles, which in turn contain subunits that either recognize appropriate polyubiquitinated targets or regulate the catalytic activity of the 20S core (Wang et al., 2005). Significantly, dysfunctions in the UPS are associated with neurodegenerative diseases such as Alzheimer's (Oddo, 2008) and Parkinson's disease (Betarbet et al., 2005). For example, one study demonstrated a decrease in proteasome activity in the

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hippocampus of Alzheimer's disease brains compared to controls (Keller et al., 2000). To further illustrate, 20S proteasome enzymatic activity is reduced in the substantia nigra pars compacta in subjects with sporadic Parkinson's disease (McNaught et al., 2003).

Our laboratory has argued for the importance of the UPS for the regulation of synaptic plasticity and memory. For example, UPS protein degradation is critical for the formation and stability of fear memories within the basolateral amygdala (BLA) (Jarome et al., 2011). Furthermore, memory formation for trace fear conditioning (TFC) requires UPS-mediated protein degradation within the pre-limbic (PL) subdivision of the medial prefrontal cortex (mPFC) (Reis et al., 2013). TFC is a variation on standard delay fear conditioning in which a stimulus-free interval is inserted between the conditioned stimulus (CS) termination and onset of the unconditioned stimulus (UCS). TFC is also unique because it engages a brain network distinct from delay conditioning. For instance, TFC requires the dorsal hippocampus (DH) and mPFC, while delay conditioning with the same stimuli does not (Esclassan et al., 2009; Gilmartin and Helmstetter, 2010). Interestingly, while delay fear conditioning remains intact throughout the lifespan, aged rats show deficits in trace memory during testing (Moyer and Brown, 2006).

In the present study, we examined whether age-related deficits in TFC memory are associated with changes in plasticity-related protein degradation processes, particularly in regard to the UPS. To these ends, we quantified the relative phosphorylation of Rpt6 and accumulation of lysine-48 (K48)-linked polyubiquitinated proteins. Rpt6 is an ATPase subunit in the 19S regulatory particle of the proteasome and is phosphorylated at Serine-120 (S120) by kinase Ca^{2+} /calmodulin-dependent protein kinase II α (CaMKII α) (Djakovic et al., 2009). In addition, CaMKII-dependent phosphorylation of Rpt6 at S120 regulates synaptic strength in hippocampal neurons (Djakovic et al., 2012). Research from our laboratory demonstrates that CaMKII regulates Rpt6 phosphorylation and proteasome activity during the formation of a long-term fear memory (Jarome et al., 2013), while inhibition of CaMKII prevents retrieval-induced increases in proteasome activity and Rpt6 phosphorylation in the amygdala (Jarome et al., 2016). We also quantified K48 polyubiquitination. There are several lysine sites on ubiquitin in which polyubiquitin chains can form, but lysine-48 (K48) linkage is only associated with the proteasome. Thus, K48 polyubiquitin chains are degradation specific and serve as an indicator of ubiquitin tagging and protein turnover. In addition, K48 polyubiquitination increases after memory retrieval (Jarome et al., 2013, 2016). Further research demonstrates that K48 specifically increases in the cytoplasmic and nuclear components of the cell during context fear conditioning consolidation, while it increases in the synaptic region during reconsolidation (Orsi et al., 2019). To understand if protein degradation processes are an important component of age-related memory impairments, we need more information about how such age-related alterations relate to these established mechanisms. In the present study, we sought to determine if age-related changes in UPS function within key brain areas known to be important for memory would reflect predicted behavioral impairments. If UPS function is specifically compromised as a function of age within one or more of the brain regions previously implicated in the retrieval of fear memory, we would expect to see a decrement in behavioral performance as a result.

2. Methods

2.1. Animals and housing conditions

Subjects were male Fisher 344 (F344) rats obtained from the National Institute on Aging colony at Charles River (Raleigh, NC,

USA) at the ages of 3, 15, and 22 months old at the time of delivery. Rats were individually housed with *ad libitum* access to water and rat chow. The animal colony was maintained at a 14:10-h light–dark cycle, with all experiments occurring under the light portion of the cycle. All experiments were approved by the Institutional Animal Care and Use Committee at the University of Wisconsin-Milwaukee and conducted within the ethical guidelines of the National Institutes of Health.

2.2. Conditioning apparatus

Fear conditioning was conducted in a set of 4 Plexiglas and stainless steel chambers within sound-attenuating boxes, as described previously (Ferrara et al., 2017). Briefly, the floor included 18 stainless steel bars connected to a shock generator (Coulbourn Instruments). Each chamber had a speaker to allow delivery of white noise cues, overhead illumination with a 7.5 W bulb, and ventilation fans to provide a constant background noise (55 dB). The chambers were cleaned with 5% ammonium hydroxide solution between sets of rats (context A). A set of similar chambers (context B) served as a shifted context for auditory CS testing. There were several distinct features associated with context B, including textured Plexiglas flooring, infrared illumination, and 5% acetic acid cleaning solution.

2.3. Trace fear conditioning procedures

All animals were handled for 3 days before behavioral manipulation. This consisted of transport to the behavior room and gentle restraint in a towel. TFC training was conducted in context A, while auditory CS testing (retrieval) was conducted in context B. All animals were trained in TFC on day 1 with 10 CS-UCS pairings after a 2-minute baseline (BL). The CS was a 10-second white noise cue (72 dB), and the UCS was a 1-second electric footshock (1 mA). The CS and UCS were separated by a 30-second trace interval (TI), and CS-UCS pairings were separated by a variable intertrial interval (ITI) of 5.25 ± 0.5 minutes. One day after conditioning, rats received a long-term memory test consisting of 2 15-second CS presentations following a 2-minute BL period. The 2 CSs were separated by an ITI of 175 seconds, and the second CS was followed by a 2-minute post-CS period. This ITI and post-CS period were combined for behavioral analysis and are referred to as the stimulus-free period (SFP) during retrieval testing (Gilmartin et al., 2012; Kwapis et al., 2011).

2.4. Behavioral scoring

Freezing behavior is defined as the cessation of all movement excluding respiration (Fanselow, 1980) and was automatically scored in real time with FreezeScan 1.0 detection software (Clever Sys, Inc) calibrated to a trained human observer.

2.5. Crude synaptosomal membrane fractionation

Animals were sacrificed with an overdose of isoflurane 90 minutes after memory testing. Brains were rapidly removed and flash frozen on dry ice. Using a rat brain matrix (Harvard Apparatus) on dry ice, the BLA, mPFC, and DH were dissected from the brains. Synaptosomal membrane fractions were obtained using methods previously described (Jarome et al., 2011) with minor alterations noted below. Tissue samples were homogenized in TEVP buffer with 320 mM sucrose and centrifuged at $1000 \times g$ for 10 minutes at 4 °C. The supernatant was collected and spun at $10,000 \times g$ for 10 minutes at 4 °C. The resulting pellet containing the synaptosomal fraction was resuspended in phospho-homogenization buffer (50 mM Tris-HCl, 6 mM sodium deoxycholate, 150 mM NaCl, 1 mM

NaF, 2 mini EDTA-free complete protease inhibitor tablets (Roche), 0.1% SDS, 1 mM sodium orthovanadate) and measured using a 660 nm protein assay (Pierce).

2.6. Western blot method

After synaptosomal preparation, protein levels were normalized and were loaded into a 7.5% sodium dodecyl sulfate–polyacrylamide gel electrophoresis gel and then transferred to polyvinylidene fluoride membranes using a Turbo Transfer System (BioRad). Membranes were incubated in 5% milk in Tris-buffered saline (TBS) + 0.1% Tween-20 (blocking buffer) for 1 hour before being incubated in primary antibody solutions for phosphorylated Rpt6–Serine120 (pRpt6; ProSci, 1:850), total Rpt6 (tRpt6; Enzo Life Sciences, 1:825), K48 polyubiquitin (Cell Signaling, 1:500), or actin (Cell Signaling, 1:1000) and 3% bovine serum albumen in TBS + 0.1% Tween-20 overnight at 4 °C. Membranes were then incubated in the appropriate secondary antibody (1:20,000) in blocking buffer for 60 minutes. Following a final wash, membranes were prepped in a chemiluminescence solution for 3 minutes. Images were captured and densitometry performed using National Institutes of Health Genesys software. The phosphorylated Rpt6–Serine120 rabbit polyclonal antibody was generated commercially (ProSci) against a synthetic peptide [NH₂-CALRND(pS)YTLHK-OH] as described previously (Djakovic et al., 2012; Jarome et al., 2013).

2.7. Data analysis

A two-way repeated-measures analysis of variance (ANOVA) was used to compare mean percent time freezing during training, and a one-way ANOVA was used to examine the change from baseline freezing during the SFP and CS periods during the retrieval test session. These were followed by Fisher's least significant differences post hoc tests. One-way ANOVAs followed by Fisher's least significant differences post hoc tests were also used to compare mean optical density across groups for proteins of interest using western blots. Normalized western blot samples are expressed as a percentage of TFC-trained 3-month-old animals. Statistical outliers were screened according to the methods outlined in the study by Field (2005). The data presented in this article excludes, in total, one outlier from the 22-month-old condition. This 22-month-old animal was an outlier on 2 measures, BLA-pRpt6 ($Z = 2.345$) and BLA-K48 ($Z = 2.078$). Thus, this one animal was removed from all subsequent analyses, resulting in final sample sizes of $n = 10$, 9, and 9 (3-, 15- and 22-month-old animals, respectively). Statistical significance was defined as $p < 0.05$. Data are presented as mean \pm standard error of the mean.

3. Results

3.1. Aging results in deficits in trace fear conditioning

Previous work has demonstrated that aged rats have deficits in TFC (Moyer and Brown, 2006). In the present study, we aimed to replicate and extend these results. We first found that 3-, 15-, and 22-month-old F344 rats perform equivalently during training (Fig. 1A) showing normal reactions to footshock. Specifically, we found a significant main effect of time ($F_{(1, 25)} = 1059$, $p < 0.0001$), a trend toward a main effect of age ($F_{(2, 25)} = 2.824$, $p = 0.0784$) and no significant interaction ($F_{(2, 25)} = 0.3279$, $p = 0.7235$). To test the long-term retention of fear memory, animals received a retrieval session in a shifted context 24 hours after training. We quantified behavioral performance at retrieval by calculating the difference between each animal's baseline freezing at retrieval and the mean of that animal's SFP freezing (i.e., the ITI and post-CS period, 295 seconds in total) to better account for individual differences in baseline values (Lattal, 1999). Here, we found a significant overall effect of age ($F_{(2, 25)} = 3.615$, $p = 0.0418$). Importantly, in post hoc comparisons, the 3- and 15-month-old animals froze more during the SFP than 22-month-old animals ($p = 0.0142$ and $p = 0.0831$, respectively; Fig. 1B). SFP freezing in TFC-trained rats reflects a conditional response to the offset itself and thus can be considered a conditional response to the CS (Gilmartin et al., 2012). However, as a secondary measure, the CS freezing was also analyzed as a difference from the baseline freezing (mean percent CS freezing–mean percent BL freezing), but we did not observe any significant differences ($F_{(2, 25)} = 0.5827$, $p = 0.5658$). Together, these results suggest that aging results in a deficit in long-term memory for TFC, specifically when one looks at the SFP period.

3.2. Aging is associated with changes in protein degradation processes

Next, we identified how memory-driven UPS signaling changes with age in several brain regions critical for TFC, including the BLA, mPFC, and DH. Prior work shows that the acquisition or retrieval of fear memory increases UPS-related signaling at amygdala synapses (Jarome et al., 2011; Orsi et al., 2019). Through western blot analysis of BLA tissue (Fig. 2A), we quantified the phosphorylation of proteasome regulatory subunit Rpt6 at Serine-120, a site known to be critical for the regulation of increases in proteasome activity and activity-dependent changes in synaptic strength (Djakovic et al., 2012). We found that age was associated with a significant difference in phosphorylated Rpt6 (pRpt6) protein expression (Fig. 2B; $F_{(2, 25)} = 4.024$, $p = 0.0305$). More specifically, 3- and 15-month-old

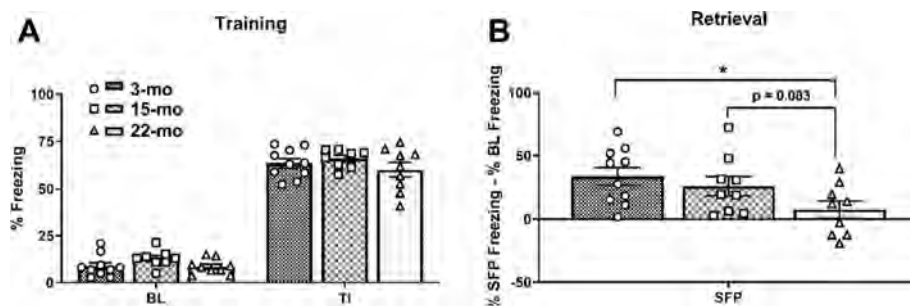


Fig. 1. Aging results in deficits in trace fear conditioning memory retrieval. (A) During training, all animals (3- [$n = 10$], 15- [$n = 9$], and 22-month-old rats [$n = 9$]) froze equivalently at BL and during the TIs. (B) However, during retrieval testing 24 hours later, significant age-related deficits in long-term memory were noted. Specifically, the SFP period, 22-month-old rats froze significantly less than 3-month-old rats and tended to freeze less than 15-month-old rats. Data are presented as mean \pm standard error of the mean. *Denotes $p < 0.05$ in a Fisher's LSD post hoc test. Abbreviations: BL, baseline; LSD, least significant differences; SFP, stimulus-free period; TI, trace interval.

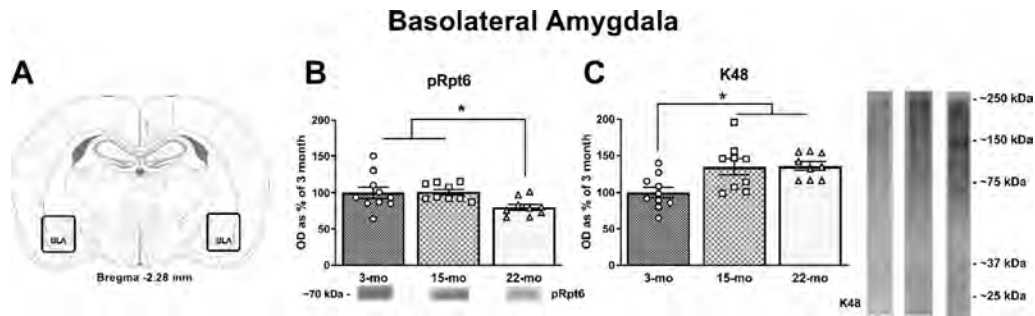


Fig. 2. Aging is associated with changes in markers of protein degradation in the amygdala. (A) Tissue was collected from the BLA (adapted from Paxinos and Watson, 2007). (B) 22-month-old rats displayed deficits in pRpt6 protein levels compared to 3- and 15-month-old animals. (C) 15- and 22-month-old rats also displayed significant increases in K48 compared to 3-month-old animals. Data are presented as mean ± standard error of the mean. *Denotes $p < 0.05$ in a Fisher's LSD post hoc test. Abbreviations: LSD, least significant differences.

animals displayed higher levels of pRpt6 compared to 22-month-old animals ($p = 0.0218$ and $p = 0.0199$, respectively). However, tRpt6 did not differ across age groups ($F_{(2, 25)} = 0.7646$, $p = 0.4761$), supporting the lack of an age-related difference in the total number of proteasomes present. We also performed western blots for a lysine-48 (K48) polyubiquitin tag that targets proteins for degradation by the proteasome (Jarome et al., 2013, 2016). We found that K48 protein levels were significantly altered by age (Fig. 2C; $F_{(2, 25)} = 6.599$, $p = 0.0050$). Specifically, both 15- and 22-month-old rats had significantly higher K48 protein levels compared to 3-month-old rats ($p = 0.0052$ and $p = 0.0040$, respectively). Finally, actin levels displayed no significant differences ($F_{(2, 25)} = 0.8937$, $p = 0.4218$). Altogether these data suggest that activity-dependent protein degradation processes are impaired in the BLA, which leads to an accumulation of ubiquitin-tagged proteins.

While plasticity in the BLA appears to be generally important for aversive learning, neurons in the mPFC and DH are specifically required when a trace interval intervenes between the CS and UCS (Escassan et al., 2009; Gilmartin and Helmstetter, 2010). Research from our laboratory has also demonstrated a functional role for prefrontal UPS-mediated degradation in the consolidation of a TFC memory (Reis et al., 2013). Aging has also affects mPFC neuronal excitability and hinders the extinction of a TFC memory (Kaczorowski et al., 2012). Overall, these results suggest that the mPFC may likely show age-related declines in UPS activity; thus, we quantified the same proteins (as in the BLA) in tissue samples collected from the mPFC (Fig. 3A). We found no significant effect of pRpt6 (Fig. 3B; $F_{(2, 25)} = 2.951$, $p = 0.0707$) and a tendency toward an effect of tRpt6 ($F_{(2, 25)} = 0.0709$, $p = 0.9318$). We also analyzed K48 protein levels in the mPFC (Fig. 3C; $F_{(2, 25)} = 6.123$, $p = 0.0068$) and found that 22-month-old animals display a tendency toward higher

levels of K48 polyubiquitination compared to 3-month-old rats ($p = 0.0654$) and significantly higher levels compared to 15-month-old rats ($p = 0.0018$). Actin, on the other hand, displayed no significant differences ($F_{(2, 25)} = 1.27$, $p = 0.2984$).

The DH is another brain region that is critical for learning the association between a CS and a UCS when a trace interval is interposed between the 2 stimuli; for instance, lesions of the DH impair trace eyeblink conditioning (Weiss et al., 1999). In another study, increases in DH CA1 pyramidal cell firing to the CS and UCS were diminished in aged animals that were unable to learn trace eyeblink conditioning, while these aged, learning-impaired animals also showed alterations in the coordinated firing of all excitatory and inhibitory pyramidal cell types within CA1 ensembles (McEchron et al., 2001). Intra-DH CA1 infusions of a proteasome inhibitor also blocks long-term memory formation following inhibitory avoidance training (Lopez-Salon et al., 2001). Therefore, we also used western blots to analyze tissue from the DH (Fig. 4A) for markers of UPS activity in young and aged animals. No differences were noted in pRpt6 (Fig. 4B; $F_{(2, 25)} = 1.164$, $p = 0.3285$), while a trend toward a difference in tRpt6 was noted ($F_{(2, 25)} = 3.02$, $p = 0.0669$). K48, on the other hand, did differ as a function of age (Fig. 4C; $F_{(2, 25)} = 17.72$, $p < 0.0001$), and post hoc analyses revealed that 22-month-old animals had significantly higher protein levels compared to 3- and 15-month-old animals (p 's < 0.0001). Actin, again, did not differ across age groups ($F_{(2, 25)} = 2.248$, $p = 0.1265$).

To better understand the role of phosphorylation of Rpt6 in TFC memory retrieval, we divided 15- and 22-month-old animals into "impaired" and "unimpaired" subgroups, as is commonly done in the field of aging and memory (Gallagher et al., 2015; Gazzaley et al., 2005; Le Jeune et al., 1996). We performed a median split using the percent time spent freezing during the SFP period at

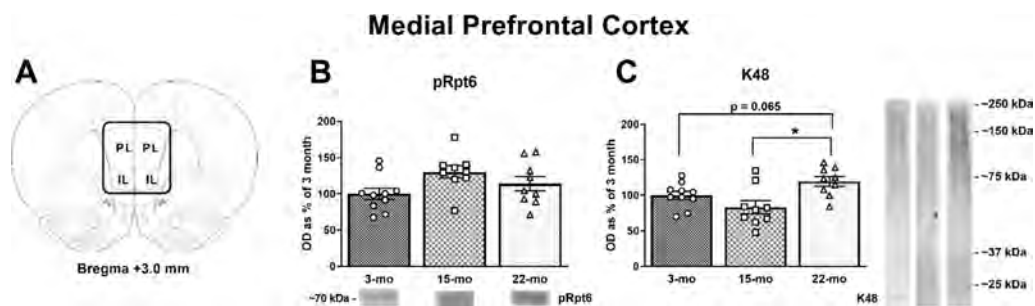


Fig. 3. Aging is associated with increases in K48 in the medial prefrontal cortex. (A) Tissue was collected from the mPFC (adapted from Paxinos and Watson, 2007). (B) No significant differences in pRpt6 were noted. (C) 22-month-old animals displayed a tendency toward increased levels of K48 compared to 3-month-old animals, significant increases in K48 compared to 15-month-old animals. Data are presented as mean ± standard error of the mean. *Denotes $p < 0.05$ in a Fisher's LSD post hoc test. Abbreviations: LSD, least significant differences.

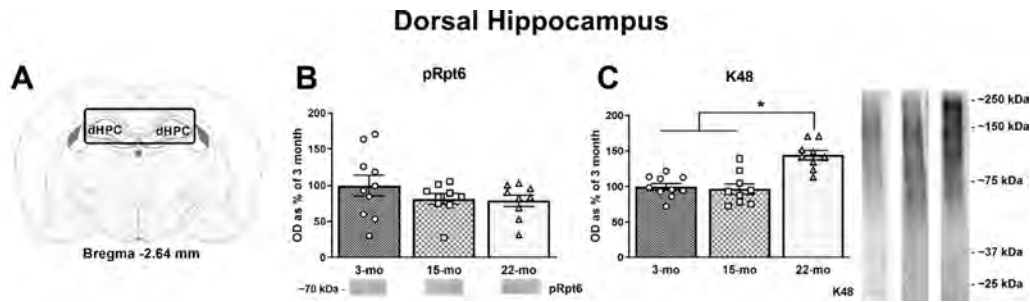


Fig. 4. Aging is associated with increases in K48 in the dorsal hippocampus. (A) Tissue was collected from the DH (adapted from Paxinos and Watson, 2007). (B) No differences were observed in pRpt6. (C) 22-month-old rats displayed significant increases in K48/Actin compared to 3- and 15-month-old rats. Data are presented as mean \pm standard error of the mean. *Denotes $p < 0.05$ in a Fisher's LSD post hoc test. Abbreviations: LSD, least significant differences.

retrieval testing (57.08%) and classified those above the median as unimpaired and those below as impaired. We then analyzed pRpt6 protein expression in each brain region in relation to the presence of behavioral impairment (Fig. 5). Not surprisingly, during the SFP of the retrieval session, impaired animals froze significantly less than unimpaired animals ($t(16) = 5.842, p < 0.0001$). Importantly, in the BLA, impaired animals had significantly lower pRpt6 protein levels ($t(16) = 2.167, p = 0.0457$). We did not find any significant differences in the mPFC ($t(16) = 0.576, p = 0.5727$) or DH ($t(16) = 0.7723, p = 0.4512$) using this analysis.

4. Discussion

Here, we provide new data supporting the idea that age-related memory deficits are associated with changes in brain protein degradation processes. This work is particularly novel and important because little research within the field of aging has focused on the amygdala as a potentially critical brain region underlying age-related memory deficits. We found that aging resulted in significant deficits in TFC memory retrieval and in changes in the expression of proteins associated with the UPS. Phosphorylation of Rpt6 at S120 is required for CaMKII α -dependent stimulation of the proteasome (Djakovic et al., 2009, 2012), and we found that normal aging is associated with decreases in pRpt6 in the BLA. We also observed an increase in K48 linked protein in aged animals in the BLA, mPFC, and DH. This increase in polyubiquitinated protein suggests that there is an accumulation of tagged protein, potentially as a result of deficits in proteasome activity (Kim et al., 2011). Taken together these data suggest that aging results in deficits in UPS function which may be closely related to TFC memory retrieval impairments.

In the present study, we chose to focus on TFC memory retrieval. Rats were sacrificed 90 minutes after retrieval since this time point shows maximal activation of the UPS after auditory fear memory retrieval (Jarome et al., 2011). However, within the dorsal hippocampus, previous studies have shown that maximal pRpt6 protein levels are sometimes observed within 0–30 minutes following context exposure (Cullen et al., 2017), and maximal polyubiquitination was observed 60 minutes following the retrieval of a context fear memory (Lee et al., 2008). Thus, our selection of a 90-minute timepoint may have missed the maximal age-related differences in protein degradation processes in some brain regions. Another potential limitation in the present study is that we are unable to determine whether these differences are dependent on the act of memory retrieval per se, or if they reflect stable age-related baseline changes in UPS function. However, in one study of the aging brain, no significant differences were noted in baseline 20S proteasome activity within the hippocampus of young and old rats (Giannini et al., 2013). On the other hand, in another study, the basal level of polyubiquitinated proteins in the hippocampus was increased with age (Zeier et al., 2011). Thus, additional research would be necessary to determine baseline levels of Rpt6 phosphorylation across several brain regions.

Another caveat is that it is difficult to delineate whether aged rats fail to initially learn TFC or if they fail to consolidate the memory. During training, equivalent freezing across groups during the TI period seems to indicate that the animals learned something about the relationship between the CS and the trace interval and respond normally to shock. TI freezing at training can be compared to SFP freezing at retrieval. However, 24 hours after training when animals were tested for the long-term retention of that fear memory, aged animals froze less during the SFP period than their

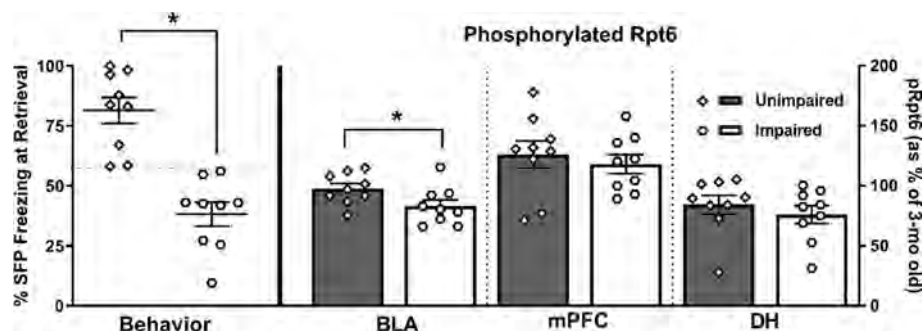


Fig. 5. Memory impaired rats display reductions in pRpt6 within the basolateral amygdala. When 15- and 22-month-old rats were classified as impaired or unimpaired, impaired animals not only display significantly less freezing during the SFP period of memory retrieval testing but also display significant reductions in pRpt6 in the BLA, but not in the mPFC or DH. Data are presented as mean \pm standard error of the mean. *Denotes $p < 0.05$ in a Student's t-test. Abbreviations: SFP, stimulus-free period.

younger counterparts. SFP freezing is commonly used to measure learning of TFC, as it represents CS-induced fear to the trace interval and encompasses the time at which the TFC-trained animal would have received the footshock during training (Blum et al., 2006; Detert et al., 2008; Kwapis et al., 2011; Quinn et al., 2002; Yoon and Otto, 2007). Finally, the present study does not provide insight into sex-specific differences in protein degradation processes. Given the gender differences in age-related neurodegenerative conditions such as Alzheimer's disease (Mielke et al., 2014) it is important that future studies directly consider sex as a biological variable.

While we observed a significant reduction in pRpt6 in the BLA of aged animals with a corresponding increase in K48 in the same region, we did not observe reductions in pRpt6 in the mPFC or DH (where increases in K48 were also noted). Although we analyzed the biochemistry of the BLA, mPFC, and DH independently, it is important to keep in mind that these brain regions are likely acting as a circuit. Differences in pRpt6 in the BLA may have functional influences on K48 in other regions via a larger circuit interaction is consistent with a growing body of research that is characterizing TFC-related circuits. For example, we have previously demonstrated that functional disconnection of the PL with the BLA impairs the acquisition of TFC (Gilmartin et al., 2012). TFC memory formation is also prevented when the mPFC is silenced specifically during the TI using optogenetic inhibition (Gilmartin et al., 2013). However, circuit-related issues in aging remain relatively unexplored. This is an important area for future research.

Proteolytic signaling pathways are critical in synapse development, synaptic plasticity, and the maintenance of neuronal health (for reviews see Bingol and Sheng, 2011; Jarome and Helmstetter, 2013). Once activated, the UPS can potentially regulate a large number of downstream signaling pathways and synaptic plasticity processes. For instance, the transcription factor cAMP response element binding protein is a downstream effector of proteasome activity (Ehlers, 2003). Overall, Ehlers (2003) demonstrated that activity regulates postsynaptic composition and signaling through the UPS, which may serve as a link between synaptic activity, protein turnover, and the reorganization of synapses. The results of the present study suggest that plasticity processes tied to the UPS change with age. UPS activity and protein clearance is impaired in the amygdala as observed through decreased phosphorylation of Rpt6 and increased K48 polyubiquitination. This aggregation of proteins in aged rats is further observed in the mPFC and DH. Importantly, when animals are classified as impaired or unimpaired based on behavioral performance, pRpt6 protein levels are correlated with the degree of impairment. One possible mechanism that may explain this correlation is a decrease in CaMKII activation in learning-impaired animals, as we have previously shown that CaMKII regulates the phosphorylation of Rpt6 and promotes memory destabilization following retrieval (Jarome et al., 2016). We have also shown that CaMKII regulates pRpt6 during the formation of a long-term memory (Jarome et al., 2013). Additionally, there is a growing body of evidence for a role for N-methyl-D-aspartate receptor in this CaMKII-UPS pathway (Jarome et al., 2011, 2013). Furthermore, N-methyl-D-aspartate receptor dysfunction has been linked to age-related oxidative stress. For instance, this age-related, oxidative stress-linked NMDA receptor hypoactivation has been observed in the hippocampus (Kumar and Foster, 2013), mPFC (Guidi et al., 2015), and amygdala (Zhan et al., 2018). Altogether, these changes in UPS-associated proteins may be responsible for age-related deficits in TFC memory, and these proteins could potentially serve as novel targets for the treatment and prevention of age-related cognitive decline.

5. Conclusions

In short, the present study provides new evidence that normal aging is associated with deficits in brain protein degradation processes. After confirming that aging results in significant deficits in TFC memory (see Moyer and Brown, 2006), we found that activity-dependent protein degradation processes are impaired, most notably in the BLA. Across several brain regions (BLA, mPFC, and DH), we also observed increases in accumulated proteins tagged for degradation with K48 polyubiquitination. Taken together, these data provide compelling evidence that aging results in changes in quality control protein degradation processes, specifically with regard to the UPS. Future research aimed at investigating methods of rescuing proteasome activity to reinstate normal learning and memory in aged animals will have important implications for the treatment and prevention of cognitive decline and neurodegenerative pathologies such as Alzheimer's disease.

Disclosure statement

The authors declare no competing financial interests.

CRediT authorship contribution statement

Brooke N. Dulka: Methodology, Investigation, Formal analysis, Writing - review & editing. **Shane E. Pullins:** Methodology, Investigation. **Patrick K. Cullen:** Investigation. **James R. Moyer:** Methodology, Writing - review & editing. **Fred J. Helmstetter:** Methodology, Formal analysis, Writing - review & editing.

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