

# Females, but not males, require protein degradation in the hippocampus for contextual fear memory formation

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Strong evidence supports a role for protein degradation in fear memory formation. However, these data have been largely done in only male animals. Here, we found that following contextual fear conditioning, females, but not males, had increased levels of proteasome activity and K48 polyubiquitin protein targeting in the dorsal hippocampus, the latter of which occurred at chaperones or RNA processing proteins. In vivo CRISPR–dCas9-mediated repression of protein degradation in the dorsal hippocampus impaired contextual fear memory in females, but not males. These results suggest a sex-specific role for protein degradation in the hippocampus during the consolidation of a contextual fear memory.

[Supplemental material is available for this article.]

The ubiquitin-proteasome system (UPS) regulates the degradation of most short-lived proteins in cells (Hershko and Ciechanover 1998) and has been widely implicated in synaptic plasticity (Ehlers 2003; Dong et al. 2008; Djakovic et al. 2012; Hegde 2017) and the pathophysiology underlying numerous neurodegenerative and psychiatric disorders (Rubio et al. 2013; Zheng et al. 2016; Cheon et al. 2019). In this pathway, target substrates get marked for degradation via the attachment of the small protein modifier ubiquitin. Multiple ubiquitin linked together at lysine-48 (K48 polyubiquitination) provides the maximal signal for degradation and these substrates are then degraded by the multisubunit 26S proteasome complex (Bedford et al. 2010). Over the last two decades strong evidence has emerged implicating UPS-mediated protein degradation in activity- and learning-dependent synaptic plasticity (Ehlers 2003; Dong et al. 2008; Jarome et al. 2011; Jarome and Helmstetter 2013; Reis et al. 2013; Rosenberg et al. 2016a,b; Hegde 2017). In the dorsal hippocampus, accumulating evidence supports a role for protein degradation in the formation or consolidation of memories for objection recognition and aversive and nonaversive spatial tasks (Lopez-Salon et al. 2001; Artinian et al. 2008; Figueiredo et al. 2015; Furini et al. 2015), and the context pre-exposure facilitation effect (Cullen et al. 2017). However, intradorsal hippocampus injections of proteasome inhibitors do not impair memory for foreground contextual fear memories (Lee et al. 2008), suggesting that some hippocampus-dependent memories do not require protein degradation during the consolidation process. Additionally, to date, the majority of the protein targets of the degradation process during any stage of memory storage remain to be identified, limiting our understanding of the functional significance for protein degradation during memory consolidation.

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Article is online at <http://www.learnmem.org/cgi/doi/10.1101/lm.053429.121>.

While the evidence supporting a role for protein degradation in memory formation is strong, a majority of it has been completed in only male animals. Recently, we reported that while both male and female rats required protein degradation for contextual fear memory formation in the amygdala, they differed in their engagement and regulation of this process following learning (Devulapalli et al. 2021). Furthermore, we found that in the amygdala and dorsal hippocampus male and female rats differed in how the proteasome could be regulated by protein kinase A (PKA) and calcium/calmodulin-dependent protein kinase II (CaMKII) following learning (Devulapalli et al. 2019). Additionally, recent evidence suggests sex differences in how protein degradation is engaged following memory retrieval in aged animals (Dulka et al. 2021). However, to date, these remain the only studies that have directly compared the protein degradation process in male and female rodents following behavioral training or retrieval. Importantly, whether males and females differ in the requirement for protein degradation, or lack thereof, in the dorsal hippocampus during the consolidation of a contextual fear memory remains unknown. Here, we directly tested this using a combination of behavioral training with biochemical assays, proteomic analyses and genetic manipulations.

Please see the **Supplemental Material** for expanded details on the methodology used. Briefly, all experiments used 8- to 9-wk-old male or female Sprague Dawley rats that were housed two per cage. For all experiments, males and females were ran at separate times, which was necessary due to the complexity of our analyses and available housing space. Animals underwent a foreground

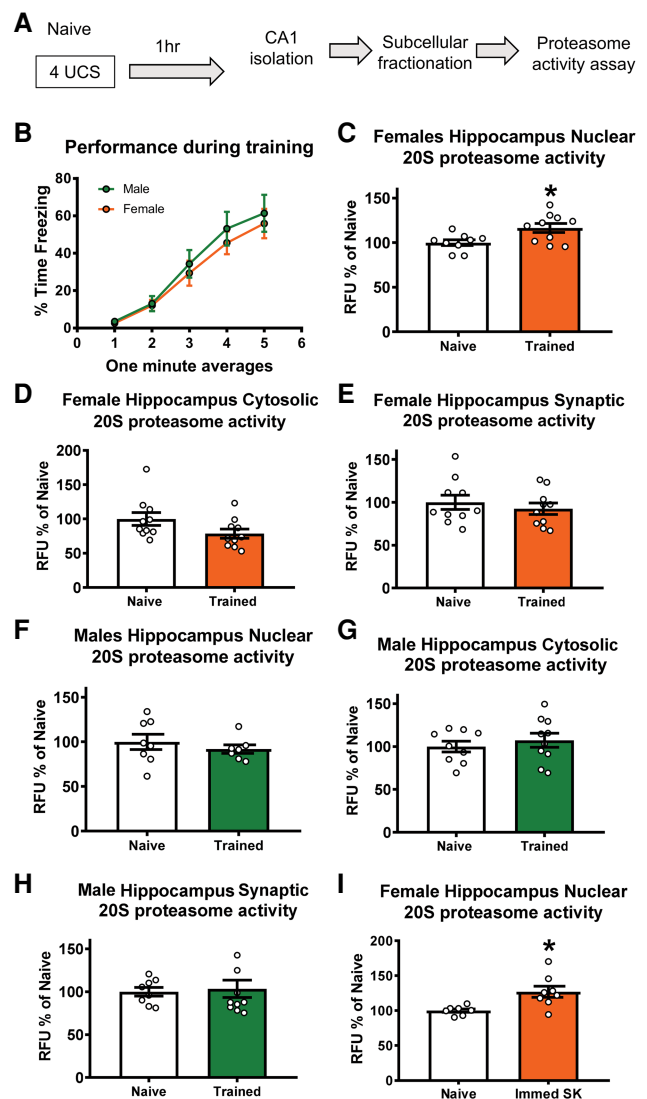
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contextual fear conditioning procedure, as previously described by our group (Orsi et al. 2019; Devulapalli et al. 2021). In some cases, the CA1 region of the dorsal hippocampus was dissected, subcellular fractions collected and proteasome activity quantified (Orsi et al. 2019; Devulapalli et al. 2021). In other cases, dorsal hippocampus tissue was dissected and purified with a K48-specific tandem ubiquitin binding entity (TUBE) followed by liquid chromatography mass spectrometry (LC/MS). Prior to behavioral training, some animals received stereotaxic injections of CRISPR-dCas9 plasmids into the CA1 region of the dorsal hippocampus using our recently described *in vivo* procedure (Devulapalli et al. 2021; Jarome et al. 2021). These animals underwent behavioral training 2 wk later and were tested for retention to the training context the following day.

We first tested whether proteasome activity was increased in the dorsal hippocampus of male and female rats 1 h after contextual fear conditioning (Fig. 1A), as this is the time point at which numerous studies have reported the earliest increase in protein degradation following fear conditioning (Jarome et al. 2011, 2013; Reis et al. 2013; Orsi et al. 2019; Devulapalli et al. 2021). Performance during the training session for both male and female rats is shown in Figure 1B. In females, we found increased proteasome activity in the nuclear ( $t_{17}=2.676$ ,  $P=0.0160$ ) (Fig. 1C), but not cytosolic ( $t_{18}=1.843$ ,  $P=0.0819$ ) (Fig. 1D) or synaptic ( $t_{18}=0.6993$ ,  $P=0.4933$ ) (Fig. 1E), fractions. Conversely, in males we did not observe changes in proteasome activity in the nuclear ( $t_{13}=0.7876$ ,  $P=0.4451$ ) (Fig. 1F), cytosolic ( $t_{17}=0.6948$ ,  $P=0.4966$ ) (Fig. 1G) or synaptic ( $t_{16}=0.2738$ ,  $P=0.7877$ ) (Fig. 1H) fractions. These results suggest that protein degradation is increased in the dorsal hippocampus of female, but not male, rats following contextual fear conditioning.

To test whether the increase in proteasome activity in females was specific to the context-shock association, we next compared naïve animals with those that underwent an immediate shock procedure in which the context and shock are presented in a nonassociative manner (Orsi et al. 2019). Interestingly, we found that immediate shock animals had increased nuclear proteasome activity relative to naïve controls ( $t_{13}=3.031$ ,  $P=0.0097$ ) (Fig. 1I). This suggests that in females, the protein degradation process is likely being increased in the dorsal hippocampus due to context exposure alone.

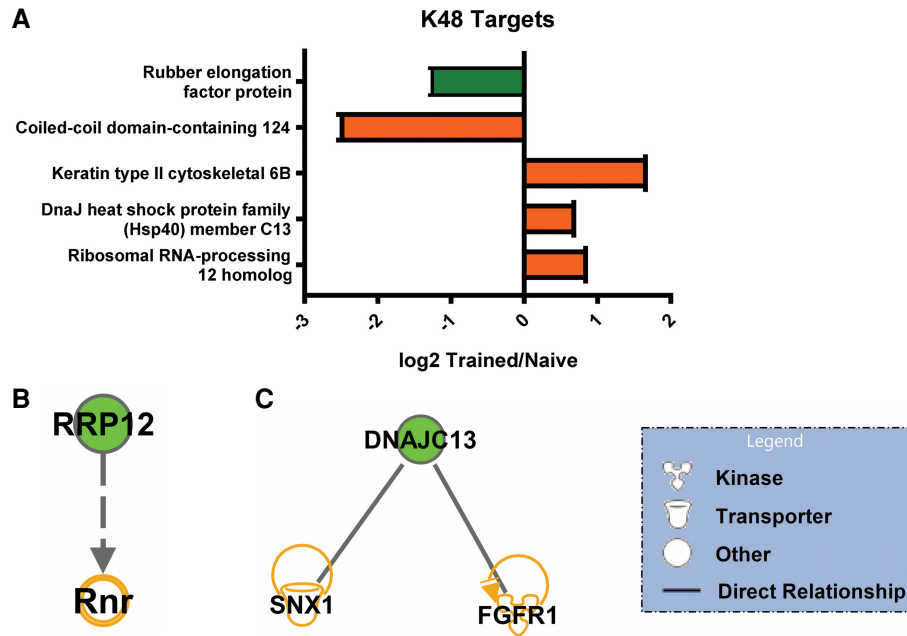
To further examine this potential sex difference in protein degradation increases in the dorsal hippocampus, we next quantified changes in degradation-specific K48 polyubiquitination in this region of male and female rats that underwent contextual fear conditioning. Broad Western blot approaches lack the sensitivity to detect changes in K48-targeting of individual proteins and we previously reported no change in global K48 polyubiquitination levels in the dorsal hippocampus of male and female rats following contextual fear conditioning (Devulapalli et al. 2021). To increase our sensitivity and identify specific targets of the protein degradation process, which could be used to infer function, we purified dorsal hippocampus tissue collected from naïve and fear conditioned male and female rats with a K48-specific TUBE followed by LC/MS. Consistent with our proteasome activity data, we detected increased K48 polyubiquitin targeting of 3 proteins in females, with 1 additional protein showing a significantly lower level of K48 targeting (Fig. 2A). Conversely, we did not identify any positive targets of K48 polyubiquitination in males, although one protein did have reduced K48 targeting as a function of learning (Fig. 2A). In females, ingenuity pathway analysis (IPA) of the protein targets of K48 polyubiquitination indicated that a downstream target of ribosomal RNA processing 12 (RRP12) is ribonucleotide reductase (RNR) (Fig. 2B), which is involved in DNA synthesis, and evidence suggests that loss of RRP12 activity is associated with a reduced function of RNR



**Figure 1.** Females, but not males, have increased proteasome activity in the dorsal hippocampus following contextual fear conditioning. (A) Experimental design showing that the dorsal CA1 region of the hippocampus was taken and dissected 1 h after contextual fear conditioning. Male and female rats were trained at separate times. (B) Behavioral performance of male and female rats during the training session. (C–E) Female rats that underwent contextual fear conditioning ( $n=10$ ) had increased proteasome activity in the nuclear (C), but not cytosolic (D) or synaptic (E), regions of cells in the hippocampus when compared with their naïve counterparts ( $n=9$  nuclear, 10 cytosolic/synaptic). (F–H) Male rats that underwent contextual fear conditioning did not have altered proteasome activity in the nucleus ( $n=7$ ; F), cytosolic ( $n=10$ ; G) or synaptic ( $n=10$ ; H) compartments relative to naïve controls ( $n=8$  nuclear/synaptic, 9 cytosolic). (I) Female rats that underwent an immediate shock procedure ( $n=8$ ) had increased proteasome activity in the nuclear compartment of cells in the hippocampus when compared with their naïve counterparts ( $n=7$ ). (\*)  $P<0.05$  from naïve.

(Tafforeau et al. 2013). Furthermore, the chaperone heat shock protein 40 (HSP40) is involved in regulation of the tyrosine kinase fibroblast growth factor receptor 1 (FGFR1) (Fig. 2C), suggesting potential functional roles of protein degradation in the DNA damage response and intracellular signaling during contextual fear memory consolidation.

Next, we tested whether males and females required protein degradation in the dorsal hippocampus for the consolidation of a

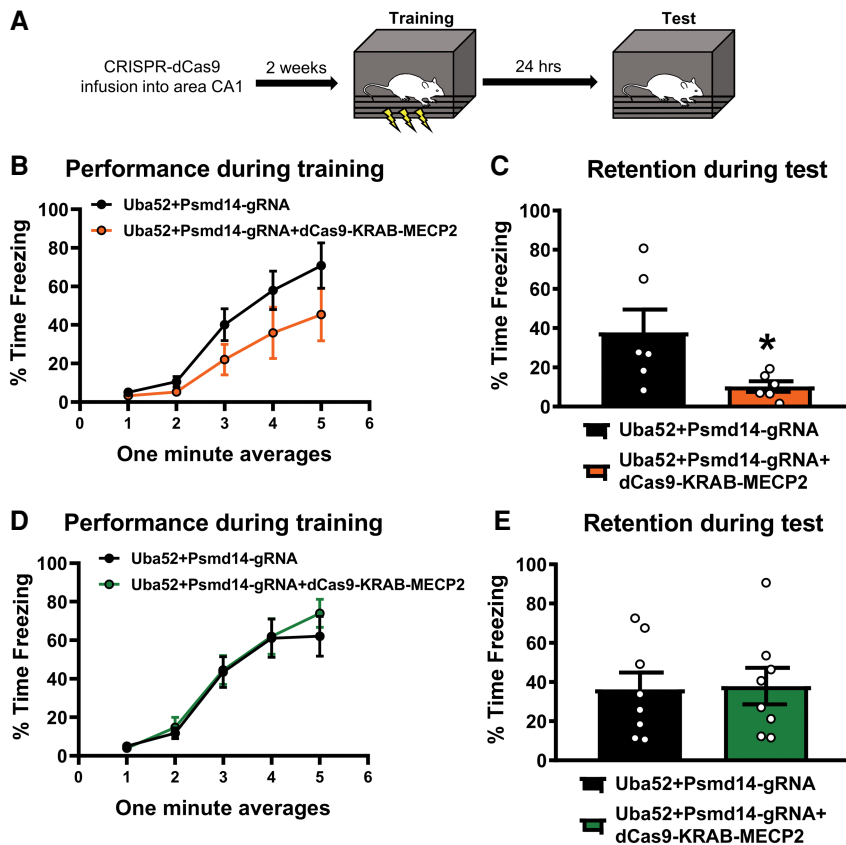


**Figure 2.** Proteomic analysis of degradation-specific K48 polyubiquitination targets in the dorsal hippocampus of male and female rats following contextual conditioning. Male and female rats were trained to a contextual fear conditioning task and the dorsal CA1 region of the hippocampus collected 1 h later for K48-specific tandem ubiquitin binding entity (K48-TUBE) liquid chromatography-mass spectrometry (MC/MS) analysis. Males and females were trained at separate times ( $n=5$  per group) and compared with same sex naive animals ( $n=5$  per group). (A) Identification of fear conditioning-induced K48 polyubiquitination targets for both females (orange bars) and males (green bar). Proteins are identified on the Y-axis and log change in K48 polyubiquitination (trained/naive) is on the X-axis. Negative log value indicates a loss of K48 polyubiquitination with fear conditioning. (B,C) Ingenuity pathway analysis (IPA) was performed on female hippocampus samples that underwent mass spectrometry analysis. (B) IPAs of downstream targets of ribosomal RNA processing 12 protein (RRP12). Green indicates a decrease in function due to K48-mediated degradation, and the dotted gray arrow denotes indirect decreased function of ribonucleotide reductase (RNR). (C) IPAs of downstream targets of heat shock protein 40 (HSP40). Green indicates a decrease in function due to K48-mediated degradation, and the solid gray arrow denotes direct inactivation of the tyrosine kinase fibroblast growth factor receptor 1 (FGFR1) and sorting nexin 1 (SNX1).

contextual fear memory. Pharmacological manipulations of the proteasome, while specific, cause an artificial accumulation of ubiquitinated proteins (Jarome et al. 2011) that can disrupt a variety of cellular processes independent of protein degradation (Pavlopoulos et al. 2011; Musaus et al. 2020), making interpretation of behavioral outcomes difficult. To overcome this, we recently developed an *in vivo* CRISPR-dCas9 (dCas9) procedure that can achieve simultaneous, bidirectional control of proteasome activity and protein ubiquitination levels in the brain (Devulapalli et al. 2021). This procedure multiplexes synthetic guide RNAs (gRNA) against proteasome subunit coding gene *Psm14* and ubiquitin coding gene *Uba52* in combination with the transcriptionally repressive dCas9-KRAB-MECP2 fusion, resulting in significant reductions in both degradation-specific K48 polyubiquitination levels and proteasome activity within 2 wk of infusion. Using this procedure (Fig. 3A), we tested whether protein degradation was necessary for the consolidation of a contextual fear memory in male and female rats. Consistent with our molecular data, in females inhibition of protein degradation did not impair performance during training (two-way ANOVA: Time =  $F_{(4,20)} = 23.22$ ,  $P < 0.0001$ ; Group =  $F_{(1,5)} = 4.179$ ,  $P = 0.0963$ ; Interaction =  $F_{(4,20)} = 1.798$ ,  $P = 0.1689$ ) (Fig. 3B) but resulted in significant behavioral deficits during the retention test ( $t_{10} = 2.31$ ,  $P = 0.0435$ ) (Fig. 3C). However, in males inhibition of protein degradation did not alter performance during training (two-way ANOVA: Time =  $F_{(4,56)} = 53.74$ ,  $P < 0.0001$ ; Group =  $F_{(1,14)} = 0.1962$ ,  $P = 0.6646$ ; Interaction =  $F_{(4,56)} = 0.4251$ ,  $P = 0.7899$ ) (Fig. 3D) or the retention test ( $t_{14} = 0.136$ ,  $P = 0.8938$ ) (Fig. 3E). Collectively, these results suggest that females, but not males, require protein degradation in the dorsal hippocampus for the formation of a contextual fear memory.

The UPS-associated protein degradation process has been strongly linked to synaptic plasticity underlying memory formation, however, much of the work previously done has been exclusively in male animals. Previous work found that administration of proteasome inhibitors into the dorsal hippocampus of male rodents does not impair a context fear memory (Lee et al. 2008), but does impair formation of a contextual memory that is acquired independent of an aversive shock stimulus (Cullen et al. 2017). Here, we found that females, but not males, had increased protein degradation in the dorsal hippocampus after training on a contextual fear conditioning task, which was likely driven by processing of the context itself. Despite this, genetic inhibition of protein degradation in the dorsal hippocampus impaired contextual fear memory in female, but not male, rats. Collectively, the results suggest that males and females differ in the requirement for protein degradation in the hippocampus for contextual fear memory formation. Importantly, in females increased proteolysis appears to be necessary to process the contextual information important for establishing the CS-UCS relationship, which is in contrast to males where this same process does not appear to be required for processing of a context-shock relationship learned in a single trial (Lee et al. 2008; Cullen et al. 2017).

Interestingly, numerous studies have reported sex differences on contextual fear conditioning tasks in rodents, often times with conflicting results (Maren et al. 1994; Wiltgen et al. 2001; Graham et al. 2009; Keiser et al. 2017; Colon et al. 2018; Devulapalli et al. 2021). While it is unknown why sex differences exist for contextual fear conditioning in rodent models or what contributes to these conflicting results, it is interesting to speculate that this could be due to differences in engagement of the protein degradation



**Figure 3.** CRISPR-dCas9-mediated down-regulation of protein degradation in the dorsal hippocampus impairs memory for a contextual fear conditioning task in females but not males. (A) Experimental design showing that the fear conditioning took place 2 wk after the dorsal hippocampus CRISPR-dCas9 injection, with testing occurring 24 h after training. Synthetic guide RNAs (gRNAs) targeted ubiquitin (*Uba52*) and proteasome subunit (*Psm14*) coding genes and were combined with a transcriptionally repressive dCas9-KRAB-MECP2 fusion. Male and female cohorts were run at separate times. (B,C) In females, injection of the CRISPR-dCas9 constructs did not impact performance during training (B) but significantly impaired retention during testing (C) relative to controls ( $n=6$  per group). (D,E) In males, injection of the CRISPR-dCas9 constructs did not impact performance during training (D) or retention during testing (E) relative to controls ( $n=8$  per group). (\*)  $P < 0.05$  from Uba52 + Psm14-gRNA.

process in the dorsal hippocampus. Future studies should aim to more precisely examine whether many of the observed sex differences in contextual fear conditioning performance are related to differences in hippocampal protein degradation in male and female rodents.

Our data show that males do not show increases in or a requirement for UPS-mediated protein degradation in the dorsal hippocampus to form a contextual fear memory, which is consistent with prior work (Lee et al. 2008). However, we extend these previous results by showing that females do show increases in protein degradation following training, which are necessary for contextual fear memory consolidation. Recently, we found that males and females differ in the learning-dependent increases and regulation of, but not requirement for, protein degradation in the amygdala during contextual fear memory formation (Devulapalli et al. 2021). Importantly, in our previous work we found that protein degradation likely does not increase in the amygdala of females following training due to high resting levels of ubiquitin-proteasome activity, which we did not observe in the dorsal hippocampus. Together, these data strongly suggest that sex differences exist in baseline levels of and/or activity-dependent increases in protein degradation across the fear circuit. Considering the important

role that other brain regions play in the consolidation of contextual fear memory, such as the prefrontal cortex and the retrosplenial cortex (Rozeske et al. 2015; Todd et al. 2017; Trask et al. 2021), future studies should examine sex differences in the requirement for protein degradation across other parts of the fear circuit.

Despite strong evidence that protein degradation is involved in the storage of various types of memories across several brain regions, to date, the target substrates of the proteasome have remained elusive. Here, we report the first unbiased proteomic analysis of degradation-specific K48 polyubiquitinated protein targets in the dorsal hippocampus following learning. Interestingly, we found that in females there were only three proteins that gained the K48 polyubiquitin mark following fear conditioning, suggesting that while necessary, the protein degradation process may not be robustly engaged in the hippocampus following learning. Importantly, our data identified downstream protein effectors that have known roles in the regulation of intracellular signaling and the DNA damage response, both of which have been implicated in memory consolidation (Johansen et al. 2011; Madabhushi et al. 2015; Li et al. 2019). While we did not directly test the role of protein degradation in regulating the downstream protein targets and processes, the data presented here are exciting as they provide the first unbiased evidence suggesting a functional role for protein degradation during memory formation. Future studies will want to use this information to directly test the functional significance of ubiquitin-proteasome-mediated protein degradation in the dorsal hippocampus during the consolidation of a contextual fear memory.

One limitation of our study was that the K48-TUBE assay could only detect one type of degradation-specific polyubiquitin chains. While K48 is the primary polyubiquitin that targets proteins for degradation (Chau et al. 1989) and generally the most abundant mark in cells, others, such as K11, can lead to proteolysis as well (Matsumoto et al. 2010). However, quantification of these other degradation chains is largely impossible in brain tissue due to technical limitations (Musaus et al. 2020). This leaves open the possibility that males do engage protein degradation in the hippocampus following contextual fear conditioning, but do so via one of the noncanonical polyubiquitin chains. While we cannot fully rule out this possibility, our data showing that inhibition of protein degradation in the dorsal hippocampus did not impair fear memory in male rats strongly suggest that they do not need this process for memory consolidation.

In conclusion, we report a novel sex difference in the role of protein degradation in contextual fear memory consolidation in the dorsal hippocampus. In combination with our previous work, these data suggest that in the hippocampus the protein degradation process is more important for fear memory formation in females than males, though both sexes have a similar need for this process in the amygdala. Considering that females are twice

as likely to develop posttraumatic stress disorder (PTSD) and Alzheimer's diseases (Christiansen and Hansen 2015; Beam et al. 2018), these results could provide important information for the understanding of sex differences in fear memory formation or age-related hippocampus-dependent memory loss.

## Acknowledgments

This work was supported by National Institutes of Health (NIH) grants MH120498, MH120569, MH122414, and MH123742 (to T.J.J.).

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Received April 14, 2021; accepted in revised form June 7, 2021.