

Sex-Specific Regulation of Fear Memory by Targeted Epigenetic Editing of *Cdk5*

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ABSTRACT

BACKGROUND: Sex differences in the expression and prevalence of trauma- and stress-related disorders have led to a growing interest in the sex-specific molecular and epigenetic mechanisms underlying these diseases. Cyclin-dependent kinase 5 (CDK5) is known to underlie both fear memory and stress behavior in male mice. Given our recent finding that targeted histone acetylation of *Cdk5* regulates stress responsivity in male mice, we hypothesized that such a mechanism may be functionally relevant in female mice as well.

METHODS: We applied epigenetic editing of *Cdk5* in the hippocampus and examined the regulation of fear memory retrieval in male and female mice. Viral expression of zinc finger proteins targeting histone acetylation to the *Cdk5* promoter was paired with a quantification of learning and memory of contextual fear conditioning, expression of CDK5, and enrichment of histone modifications of the *Cdk5* gene.

RESULTS: We found that male mice exhibit stronger long-term memory retrieval than do female mice, and this finding was associated with male-specific epigenetic activation of hippocampal *Cdk5* expression. Sex differences in behavior and epigenetic regulation of *Cdk5* occurred after long-term, but not short-term, fear memory retrieval. Finally, targeted histone acetylation of hippocampal *Cdk5* promoter attenuated fear memory retrieval and increased tau phosphorylation in female but not male mice.

CONCLUSIONS: Epigenetic editing uncovered a female-specific role of *Cdk5* activation in attenuating fear memory retrieval. This finding may be attributed to CDK5 mediated hyperphosphorylation of tau only in the female hippocampus. Sex-specific epigenetic regulation of *Cdk5* may reflect differences in the effect of CDK5 on downstream target proteins that regulate memory.

Keywords: CDK5, Epigenetics, Memory, PTSD, Sexual dimorphism, Zinc fingers

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A prominent mechanism by which cells respond to environmental stimuli is regulation of histone posttranslational modifications (hPTMs) (1,2). Evidence of such epigenetic modifications in the context of fear learning and memory is widely documented (3–5). Sexually dimorphic epigenetic gene regulation (6,7) may underlie observed sex differences in pathological memory formation associated with posttraumatic stress disorder (PTSD) (8,9) as well as a host of other neuro-pathological disorders, including depression (10–12). Many of these disorders include cognitive and anxiety symptoms that are modeled by fear conditioning and memory retrieval in rodents (6,13). However, the precise molecular mechanisms by which sex-specific epigenetic regulation of a given target gene modulates behavior is poorly understood. Targeted epigenetic editing is a novel approach to elucidate the direct causal relevance of epigenetic regulation of a given gene of interest to neuropsychiatric (14,15) and neurodevelopmental (16,17) disease.

One key gene involved in both fear memory formation and stress-related behavior is cyclin-dependent kinase 5 (*Cdk5*), whose involvement has been shown through conditional

deletion in the hippocampus (HPC) (18), striatum (19), and forebrain (20) to regulate both the expression and magnitude of fear-related memory and depressive-like phenotypes in male mice. Repeated stress in male mice is accompanied by activation of CDK5, phosphorylation of glucocorticoid receptors, increased expression of histone deacetylase 2, and reduced expression of memory-related genes in the HPC (21). We recently reported that targeted epigenetic activation of *Cdk5* in the nucleus accumbens (NAc) is sufficient to attenuate a depressive phenotype following male social defeat stress (15). Recent studies also point to a role for *Cdk5* gene expression in human depression (22) and a role for *Cdk5* gene expression in sexually dimorphic stress behavior in mice (23). We hypothesized that histone acetylation of the *Cdk5* promoter in the HPC is sufficient to regulate its expression and influence fear memory formation in both male and female mice. We systematically investigated *Cdk5* gene regulation by fear-related memory and retrieval in both sexes. Using the approach of targeted histone acetylation, we identified a novel, sexually dimorphic, epigenetic mechanism that is sufficient to attenuate fear memory retrieval specifically in female mice.

METHODS AND MATERIALS

Animals and Behavioral Paradigms

Experiments used male and female C57BL/6J mice that were 8 to 10 weeks of age. All procedures were approved by the Institutional Animal Care and Use Committee of the University of Pennsylvania. Fear conditioning was performed as described previously (24) with three 2-second foot shocks (0.4 mA), separated by a 60-second interval. Memory retrieval test was measured for 5 minutes after 1 hour (short-term memory [STM]) or 24 hours (long-term memory [LTM]). Cocaine-induced conditioned place preference (CPP) was carried out as described previously (15) for 2 days, using 15 mg/kg cocaine. Additional details are presented in the Supplement.

Tissue Collection, RNA Extraction, and Quantitative Real-Time Polymerase Chain Reaction

Mice were euthanized by cervical dislocation. Hippocampal punches were dissected and processed as described previously (15). Additional details, including quantitative real-time polymerase chain reaction primers and analysis method, are provided in the Supplement.

Quantitative Chromatin Immunoprecipitation

Quantitative chromatin immunoprecipitation was performed as described previously (14) on bilateral, 1-mm-diameter punches pooled from CA1 of two mice. A detailed protocol is provided in the Supplement.

Viral-Mediated Gene Transfer

Herpes simplex virus (HSV) expressing zinc finger proteins (ZFPs) fused to p65 were prepared as previously described (15) and as detailed in the Supplement.

Protein Extraction and Western Blotting

Protein extraction and Western blotting was carried out as described previously (14). Complete details of electrophoresis conditions and antibodies used are provided in the Supplement.

Statistics

Statistical analysis was performed using GraphPad Prism version 7 (GraphPad, San Diego, CA). Data were analyzed using two-way analysis of variance with conditioning, virus, and/or sex as factors followed by Bonferroni post hoc analysis for multiple comparisons. All data are expressed as mean \pm SEM. Results were considered statistically significant when $p < .05$. Outliers were removed using Grubbs' test ($n = 1$ outlier per cohort). Sample size was determined empirically and based on published literature. Sample size is included in the figure legends.

RESULTS

Long-term Fear Memory Retrieval Differs Between Male and Female Mice

To expand understanding of the well-established role of *Cdk5* in both learning- and stress-related behavior in male mice, we focused our attention on fear-related memory in both sexes.

We quantified STM retrieval as 1 hour and LTM retrieval as 24 hours after fear conditioning (Figure 1A). Percentage time spent freezing was quantified in fear-conditioned and non-shocked, context-only control mice (Figure 1B). In both sexes, freezing behavior increased over the course of the three-shock acquisition phase (Figure 1B), freezing was greatest during LTM retrieval, and overall freezing behavior was greater in shocked mice than in nonshocked mice. Interestingly, we found that female mice displayed a reduced magnitude of LTM retrieval compared with that of male mice. There was no sex difference in acquisition or STM retrieval.

Sex-Specific Activation of CDK5 Expression Following Long-term Fear Memory Retrieval

Given the observed sex differences in LTM retrieval and the known role of CDK5 in fear memory (18,25) and stress (15) in male mice, we next examined expression of CDK5 following acquisition and short- and long-term memory retrieval (see Figure 1A). We examined CDK5 expression CA1 of the HPC, as this region is functionally relevant to fear memory formation

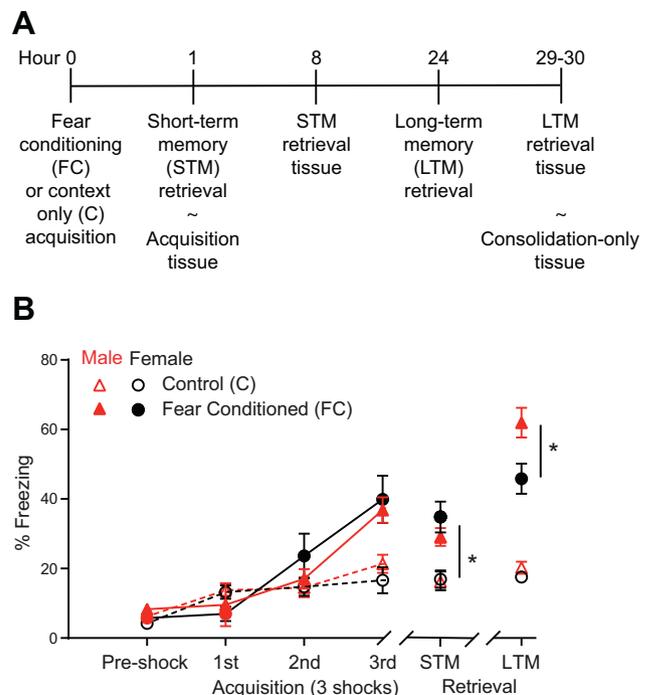


Figure 1. Long-term memory (LTM) retrieval differs between male and female mice. **(A)** Experimental timeline, in hours, depicting phases of fear conditioning and tissue collection. **(B)** Male and female mice respond differently to contextual fear conditioning, measured as percentage of time freezing (two-way analysis of variance: main effects of sex [$F_{3,35} = 9.833$, $p < .0001$], fear conditioning [$F_{4,140} = 67.67$, $p < .0001$], interaction [$F_{12,140} = 10.45$, $p < .0001$]). There was no significant difference between freezing in fear-conditioned male and female mice during acquisition (preshock, $p > .9999$; first shock, $p > .9999$; second shock, $p = .4593$; third shock, $p = .7521$) and short-term memory (STM) retrieval ($p = .2410$), but there was a lower percentage of time freezing during LTM retrieval ($p = .0377$) in female mice compared with that of male mice. In all cases, *Cdk5* expression in fear-conditioned (FC) animals was compared with that in their nonshocked, context-only control counterparts (C). All data are presented as mean \pm SEM. $n = 8-10$, * $p < .05$.

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(26–28). In order to distinguish regulation of *Cdk5* by learning and memory from that of re-exposure to the unconditioned stimuli (context), we included a consolidation-only group that underwent fear conditioning and LTM consolidation but no retrieval test.

We found that expression of *Cdk5* messenger RNA (mRNA) was increased in CA1 of male, but not female, mice after LTM retrieval, compared with its expression in the respective non-shocked, context-only control groups (Figure 2A). The male-specific increase in *Cdk5* expression was limited to the LTM retrieval phase, as there was no difference in *Cdk5* mRNA expression in CA1 of mice of either sex after acquisition or consolidation phases, compared with that found in control mice (Figure 2A). We further validated the mRNA result by Western blotting and confirmed that CDK5 protein expression in CA1 was increased after LTM retrieval in male but not female mice (Figure 2B; Supplemental Figure S1).

To determine whether fear memory regulation of CDK5 expression was specific to CA1, we next examined expression of *Cdk5* in the NAc, a reward-related brain area in which CDK5 expression is known to regulate affective behavior in male mice (15,29). There was no significant change in *Cdk5* mRNA expression after LTM retrieval in the NAc of mice of either sex (Figure 2C). We also examined mRNA expression of brain-derived neurotrophic factor from alternative promoter IV,

which is implicated in LTM retrieval (4). We found a significant increase in *Bdnf* mRNA expression following LTM retrieval in CA1 region of both sexes (Figure 2D), indicating that male-specific fear-induced gene expression is not universal.

To ensure that hormonal regulation of *Cdk5* was not responsible for the observed effects on LTM retrieval and CDK5 expression, we analyzed *Cdk5* mRNA expression in a separate cohort of naïve estrous-tracked females. We selected to compare proestrous and estrous females with males because, although proestrus lasts for short time, it shows maximum hormonal changes (30,31). No significant differences in *Cdk5* expression were measured between these groups (Supplemental Figure S2).

Sex-Specific Epigenetic Regulation of *Cdk5* Expression Following Long-term Fear Memory Retrieval

Histone acetylation and methylation of the *Cdk5* promoter activate and repress *Cdk5* expression, respectively, in brain reward regions to affect stress and depression (15,32), yet epigenetic regulation of *Cdk5* in the HPC has not yet been examined. Thus, we tested the hypothesis that histone acetylation of the *Cdk5* promoter underlies male-specific mRNA expression following fear conditioning. We used

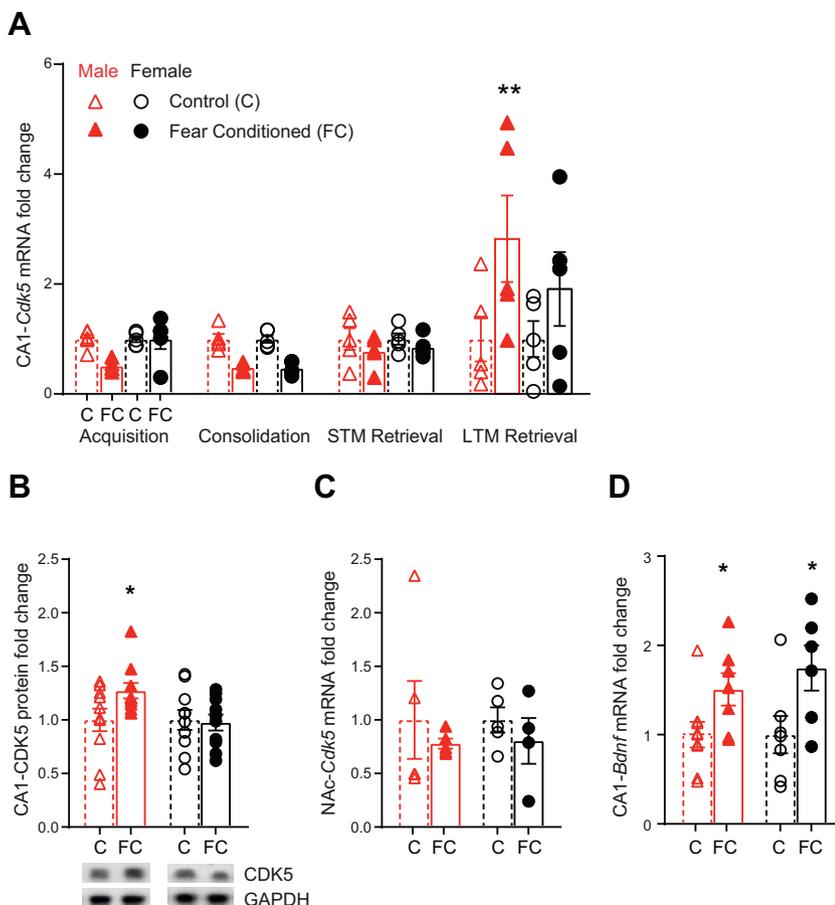


Figure 2. Sex-specific activation of cyclin-dependent kinase 5 (CDK5) expression following long-term memory (LTM) retrieval. **(A)** *Cdk5* messenger RNA (mRNA) expression was quantified by quantitative real-time polymerase chain reaction after each phase of fear conditioning (FC) (two-way analysis of variance [ANOVA], main effect of sex [$F_{1,62} = 0.04781$, $p = .8276$], and FC [$F_{7,62} = 6.666$, $p < .0001$] as factors). *Cdk5* expression was increased following LTM retrieval in CA1 of male ($p = .0096$) but not female ($p > .9999$) mice compared with that in respective behavioral control mice (C) ($n = 10$). **(B)** CDK5 protein expression in CA1 after LTM retrieval (two-way ANOVA, main effect of sex [$F_{1,36} = 5.861$, $p = .0206$], FC [$F_{1,36} = 6.266$, $p = .0170$], interaction [$F_{1,36} = 4.197$, $p = .0478$]). CDK5 protein expression was increased after LTM retrieval in male ($p = .0470$) but not in female ($p > .9999$) mice compared with that in respective behavioral control mice ($n = 10$). **(C)** *Cdk5* expression in the nucleus accumbens (NAc) did not change after LTM retrieval in mice of either sex (two-way ANOVA, main effect of sex [$F_{1,15} = 0.003143$, $p = .9560$], FC [$F_{1,15} = 0.8602$, $p = .3683$]) ($n = 4-5$). **(D)** Brain-derived neurotrophic factor (*Bdnf*) mRNA expression in CA1 was increased following LTM retrieval (two-way ANOVA, main effect of FC [$F_{1,24} = 14.33$, $p = .0009$] and sex [$F_{1,24} = 0.0006153$, $p = .9804$]) in both male ($p = .0424$) and female ($p = .0421$) mice ($n = 7-10$). All data are presented as mean \pm SEM. * $p < .05$, ** $p < .01$. GAPDH, glyceraldehyde 3-phosphate dehydrogenase; STM, short-term memory.

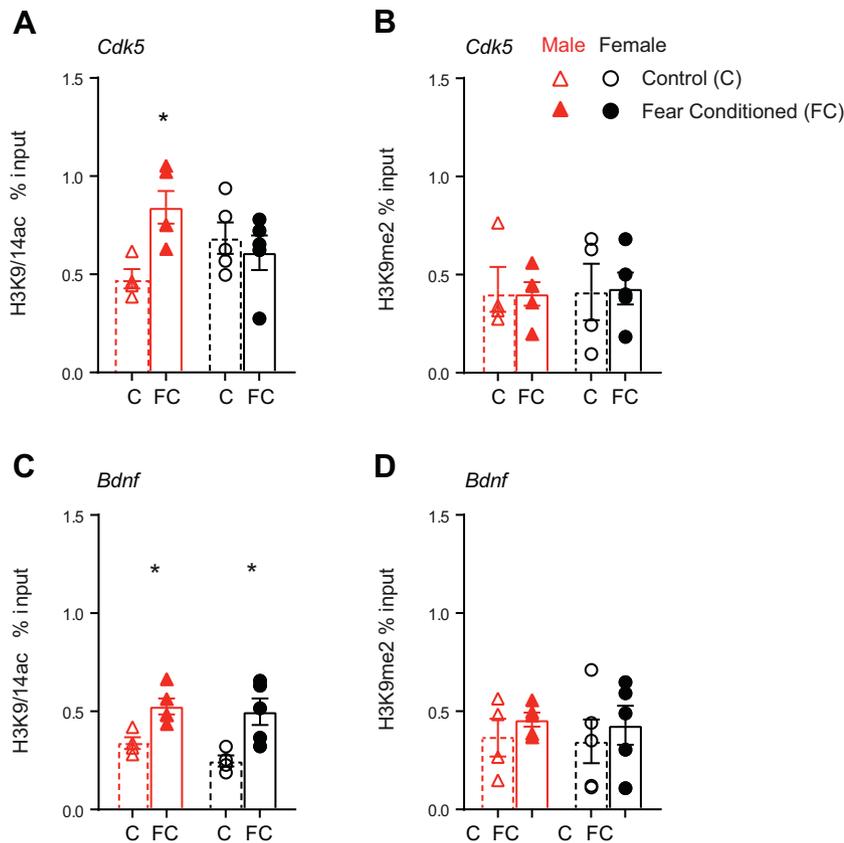


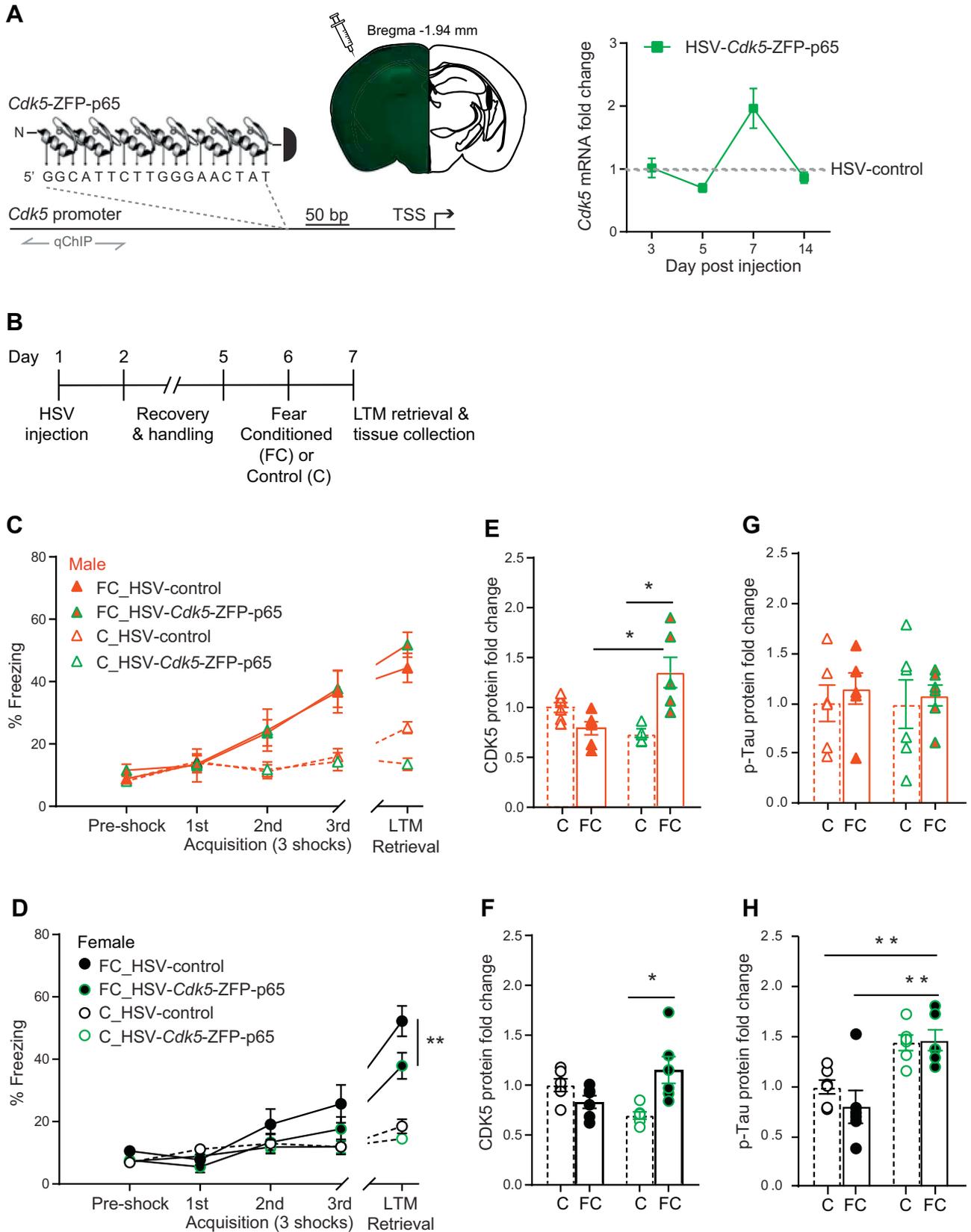
Figure 3. Sex-specific epigenetic regulation of cyclin-dependent kinase 5 (CDK5) expression following long-term memory retrieval. **(A)** Histone H3 lysine 9/14 acetylation (H3K9/14ac) enrichment of the *Cdk5* promoter in CA1 was quantified by quantitative chromatin immunoprecipitation after long-term memory retrieval (two-way analysis of variance, main effect of sex [$F_{1,15} = 0.02164, p = .8850$], fear conditioning (FC) [$F_{1,15} = 3.282, p = .0901$], interaction [$F_{1,15} = 7.504, p = .0152$]). FC enriched H3K9/14ac at the *Cdk5* promoter in CA1 of male ($p = .0415$) but not female ($p > .9999$) mice, compared with that of the respective behavioral control mice (C). **(B)** FC had no effect on enrichment of histone H3 lysine 9 dimethylation (H3K9me2) at the *Cdk5* promoter in CA1 of either male ($p > .9999$) or female ($p > .9999$) mice (two-way analysis of variance, main effect of sex [$F_{1,14} = 0.006133, p = .9387$], FC [$F_{1,14} = 0.0007834, p = .9781$], interaction [$F_{1,14} = 0.04226, p = .8401$]). **(C)** H3K9/14ac enrichment was greater at the brain-derived neurotrophic factor (*Bdnf*) IV promoter following FC (two-way analysis of variance, main effect of FC [$F_{1,14} = 23.93, p = .0002$], sex [$F_{1,14} = 2.094, p = .1699$], interaction [$F_{1,14} = 0.2557, p = .6210$]) in both male ($p = .0469$) and female ($p = .0113$) mice compared with that in respective behavioral control mice. **(D)** There was no difference in H3K9me2 enrichment at the *Bdnf* factor IV promoter following FC (two-way analysis of variance, main effect of sex [$F_{1,15} = 0.07074, p = .7939$], FC [$F_{1,15} = 0.9145, p = .3541$], interaction [$F_{1,15} = 0.003047, p = .9567$]). Quantitative chromatin immunoprecipitation data are normalized to the input of the corresponding sample. $n = 4$ or 5 chromatin immunoprecipitation samples per behavioral group. All data are presented as mean \pm SEM. * $p < .05$.

quantitative chromatin immunoprecipitation to measure enrichment of histone H3 lysine 9/14 acetylation (H3K9/14ac) and histone H3 lysine 9 di-methylation (H3K9me2), which are hPTMs associated with gene activation and repression, respectively. We examined these epigenetic changes in CA1 following LTM retrieval because of the male-specific increase

in *Cdk5* mRNA expression at this time point (see Figure 2A). H3K9/14ac was enriched at the *Cdk5* promoter in male CA1 following fear conditioning compared with that of non-shocked, context-only control mice (Figure 3A). There was no difference in H3K9/14ac at *Cdk5* in female CA1 (Figure 3A). There was no difference in *Cdk5* enrichment of

Figure 4. Epigenetic editing of the cyclin-dependent kinase 5 (*Cdk5*) promoter regulates long-term memory (LTM) retrieval in female mice only. **(A)** Schematic of zinc finger protein (ZFP)-mediated epigenetic editing and representative image of CA1-expressing herpes simplex virus (HSV)-*Cdk5*-ZFP-p65; green fluorescent protein. Time-course analysis found that maximum activation of *Cdk5* in CA1 occurred 7 days after HSV injection. *Cdk5* messenger RNA (mRNA) was analyzed by quantitative real-time polymerase chain reaction and normalized to a nonfunctional control virus, HSV-control ($n = 8$ or 9). **(B)** Timeline of mice injected with either HSV-control or HSV-*Cdk5*-ZFP-p65 and subjected to fear conditioning (FC). **(C)** Male and **(D)** female FC mice injected with either HSV-*Cdk5*-ZFP-p65 or HSV-control showed a gradual increase in freezing during acquisition (two-way analysis of variance [ANOVA], main effect of FC [$F_{4,36} = 67.92, p < .0001$], sex [$F_{3,27} = 3.946, p < .0187$], interaction [$F_{12,108} = 0.9452, p = .4969$]). Male FC mice injected with HSV-*Cdk5*-ZFP-p65 did not differ in acquisition or LTM retrieval freezing compared with that of FC mice injected with HSV-control (preshock $p > .9999$, first shock $p > .9999$, second shock $p > .9999$, third shock $p > .9999$, and LTM $p > .9999$). Female FC mice injected with HSV-*Cdk5*-ZFP-p65 displayed comparable freezing during acquisition and reduced freezing in LTM retrieval compared with that of HSV-control-injected mice (preshock $p > .9999$, first shock $p > .9999$, second shock $p > .9999$, third shock $p > .9999$, and LTM $p = .0365$). HSV-*Cdk5*-ZFP-p65-injected female mice froze less than males during LTM retrieval ($p = .0285$). Nonshocked control mice injected with HSV-*Cdk5*-ZFP-p65 did not differ from those injected with HSV-control in terms of freezing (preshock $p > .9999$, first shock $p > .9999$, second shock $p > .9999$, third shock $p > .9999$, and LTM $p > .9597$). **(E)** Male and **(F)** female Western blot analysis of CDK5 expression after LTM retrieval confirmed increased CDK5 levels in HSV-*Cdk5*-ZFP-p65-injected mice (two-way ANOVA, main effect of virus [$F_{1,5} = 9.939, p = .0253$], sex [$F_{1,5} = 1.159, p = .2213$], interaction [$F_{1,5} = 7.321, p = .0458$]). CDK5 expression in mice injected with HSV-*Cdk5*-ZFP-p65 was greater than that of HSV-control-injected mice (male $p = .0026$, female $p = .0417$). **(G, H)** Western blot analyses of phosphorylated tau protein (p-Tau) expression after LTM retrieval in HSV-*Cdk5*-ZFP-p65-injected **(G)** male and **(H)** female mice (two-way ANOVA, main effect of virus [$F_{1,5} = 16.13, p = .0102$], sex [$F_{1,5} = 10.73, p = .0221$], interaction [$F_{1,5} = 13.28, p = .0148$]). p-Tau levels in mice injected with HSV-*Cdk5*-ZFP-p65 were greater in female ($p = .0276$) but not male ($p > .9999$) mice compared with those in HSV-control-injected mice. There was no effect on p-tau levels in HSV-*Cdk5*-ZFP-p65-injected, nonshocked control mice (two-way ANOVA, main effect of virus [$F_{1,5} = 1.965, p = .2199$], sex [$F_{1,5} = 2.649, p = .1646$], interaction [$F_{1,5} = 1.351, p = .2976$]). All data are presented as mean \pm SEM. $n = 6$ –10. * $p < .05$, ** $p < .01$. bp, base pair; C, control; qChIP, quantitative chromatin immunoprecipitation; TSS, transcription start site.

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the repressive modification, H3K9me2, following LTM retrieval in mice of either sex (Figure 3B). As a positive control, we also examined histone modifications at the *Bdnf* exon IV promoter and found that H3K9/14ac was enriched in CA1 of both sexes (Figure 3C). There was no difference in the enrichment of H3K9me2 in mice of either sex at the *Bdnf* promoter (Figure 3D). These results are consistent with a male-specific increase in *Cdk5* expression and an increase in *Bdnf* expression in both sexes following fear conditioning.

Targeted Epigenetic Editing of *Cdk5* Attenuates Long-term Fear Memory Retrieval in Female Mice Only

To directly test the causal relevance of male-specific acetylation and expression of *Cdk5* in long-term fear memory consolidation, we applied targeted epigenetic editing to acetylate H3K9/14 at the *Cdk5* promoter in CA1. Engineered ZFPs were composed of six zinc fingers that uniquely bind an 18-base pair motif in the *Cdk5* promoter region (15) and regulate acetylation via fusion to the p65 transcriptional activation domain (Figure 4A). This approach mimics experience-driven transcriptional regulation of *Cdk5* in both magnitude (29,33,34) and mechanism (35,36).

Cdk5-ZFP-p65 constructs were packaged into HSV and stereotactically delivered to the CA1 region of the HPC (Figure 4A). Prior studies have found that epigenetic editing of the *Cdk5* locus regulates behaviors learned over 4 to 10 days (15). Because fear conditioning is a single-trial learning paradigm, we performed an initial time-course study and determined that the maximum activation of *Cdk5* in CA1 by HSV-*Cdk5*-ZFP-p65 occurs 7 days after HSV injection (Figure 4A). To recapitulate the timing of endogenous *Cdk5* acetylation and expression following fear conditioning (34) (see Figure 3A), we injected CA1 with HSV-*Cdk5*-ZFP-p65 and subjected mice to fear conditioning on day 6, followed by LTM retrieval on day 7 (Figure 4B). We confirmed targeting and expression of HSV-*Cdk5*-ZFP-p65 in CA1 using a fluorescent stereoscope (see representative image in Figure 4A); non-HPC-targeted or nonexpressing animals were removed from the study. We compared the effects of HSV-*Cdk5*-ZFP-p65 with those of a nonfunctional control virus, HSV-control, which expresses the p65 subunit alone (14,15).

All fear-conditioned mice increased freezing during LTM retrieval relative to that of context-only control mice (Figure 4C, D), indicating that viral injection did not interfere overall with the formation of fear memory in mice of either sex. Surprisingly, targeted acetylation of *Cdk5* had no effect on LTM in male mice (Figure 4C), while it decreased LTM in female mice, relative to respective HSV-control-injected mice (Figure 4D). Further, LTM retrieval in female HSV-*Cdk5*-ZFP-p65-injected mice was lower than that in males (compare Figure 4C, D), which is consistent with reduced freezing levels in virus-naïve female mice relative to those in male mice (see Figure 1B). There was no effect of HSV-*Cdk5*-ZFP-p65 on acquisition in mice of either sex, nor on freezing levels in nonshocked, context-only control mice (Figure 4C, D). In mice of both sexes, HSV-*Cdk5*-ZFP-p65 injection increased CDK5 expression compared with that in HSV-control-injected mice (Figure 4E, F; Supplemental Figure S3A).

We noted that in male mice injected with HSV, fear conditioning did not cause an increase in CDK5 expression (Figure 4E, F; Supplemental Figure S3) as expected based on results in virus-naïve male mice (Figure 2A). We hypothesized that surgery and anesthesia may repress *Cdk5* expression, masking the effect of fear-conditioning-activated expression in this context. This hypothesis is supported by the fact that in the rat HPC, general anesthesia decreases histone H3 acetylation and histone acetyltransferase activity of cyclic adenosine monophosphate response element binding protein-binding protein, leading to repression of brain-derived neurotrophic factor and c-Fos expression (37). To determine the effect on *Cdk5* expression of two commonly used anesthetics, ketamine and isoflurane, with and without intracranial surgery, and to recapitulate the observed effect in virus-injected fear conditioning experiments (Figure 4), we collected CA1 tissue 7 days after surgery and analyzed *Cdk5* expression by quantitative real-time polymerase chain reaction. Indeed, we found that ketamine and isoflurane anesthesia, coupled with intracranial surgery, repressed *Cdk5* expression compared with that in surgery-naïve male and female mice (Supplemental Figure S4A). We also found that HSV-*Cdk5*-ZFP-p65 expression overcomes the repression caused by anesthesia and surgery, while HSV-control does not (Supplemental Figure S4B).

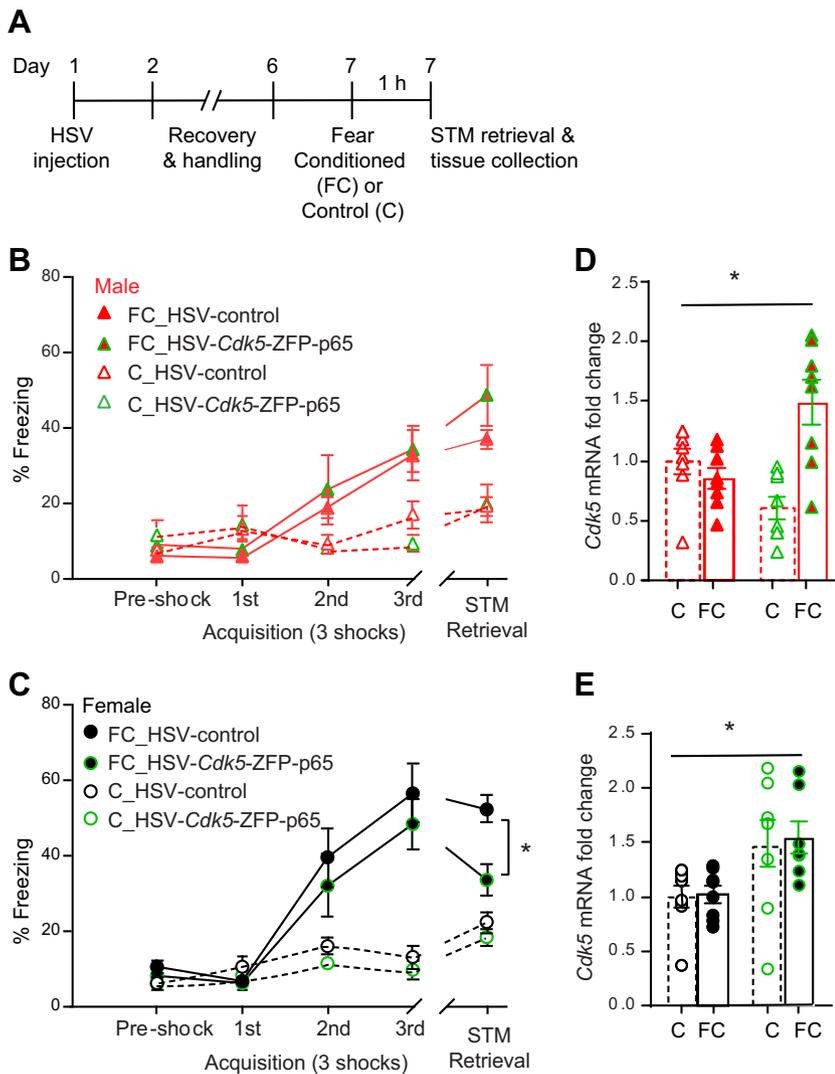
CDK5 Activation Led to Hyperphosphorylation of Tau Protein in CA1 of Female but Not Male Mice

To elucidate a potential mechanism for CDK5-mediated attenuation of LTM retrieval, we measured phosphorylation of tau protein, a direct and well-characterized downstream target of CDK5. Tau protein phosphorylation has been implicated in memory deficits, including those of working and reference memory (38,39) and spatial memory (40–42). CDK5 phosphorylates tau protein at serine 396 (43), and female-specific neurological effects of hyperphosphorylated tau protein are well documented (44–48). To determine the role of CDK5-mediated hyperphosphorylation of tau protein in memory deficits, we measured serine 396 phosphorylated tau protein relative to total tau protein in male and female mice injected with HSV-*Cdk5*-ZFP-p65 and subjected to fear conditioning. In male mice, HSV-*Cdk5*-ZFP-p65 and LTM retrieval did not change phosphorylated tau protein levels compared with those of HSV-control-injected and nonshocked, context-only male mice (Figure 4G and Supplemental Figure S3A). Conversely, in female mice, HSV-*Cdk5*-ZFP-p65 and LTM retrieval increased phosphorylated tau protein levels (Figure 4H and Supplemental Figure S3B). The female-specific phosphorylation of tau protein is consistent with female-specific attenuation of LTM retrieval following activation of *Cdk5* expression.

Targeted Epigenetic Editing of *Cdk5* Attenuates Short-term Fear Memory Retrieval in Female Mice Only and Has No Effect on Acquisition or CPP

Although *Cdk5* mRNA expression was not regulated during fear memory acquisition or STM retrieval in mice of either sex (Figure 2A), we considered that exogenous acetylation might impact these phases of learning and memory. To

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of virus [$F_{1,7} = 7.888, p = .0262$], sex [$F_{1,7} = 0.0004, p = .9839$], interaction [$F_{1,7} = 0.0003, p = .9850$]. *Cdk5* expression in mice injected with HSV-*Cdk5*-ZFP-p65 showed a trend toward increased expression compared with that of HSV-control-injected mice (male $p = .0696$, female $p = .0724$). All data are presented as mean \pm SEM. $n = 7$ or 8. * $p < .05$.

acetylate the *Cdk5* promoter during acquisition and STM retrieval, we injected CA1 with HSV-*Cdk5*-ZFP-p65 and subjected mice to fear conditioning and STM retrieval 7 days later. There was no effect on acquisition in mice of either sex (Figure 5B, C). Alternatively, while targeted *Cdk5* acetylation had no effect on STM retrieval in male mice (Figure 5B), it decreased STM retrieval in female mice, relative to that in respective HSV-control-injected mice (Figure 5C). Distinct from LTM retrieval, STM retrieval following HSV-*Cdk5*-ZFP-p65 injection was not lower in female mice than that in male mice. We measured *Cdk5* mRNA expression by quantitative real-time polymerase chain reaction 6 hours after STM retrieval. In both male and female mice, HSV-*Cdk5*-ZFP-p65 injection and STM retrieval increased *Cdk5* expression compared with that in HSV-control-injected mice (Figure 5D, E).

Figure 5. Epigenetic editing of the cyclin-dependent kinase 5 (*Cdk5*) promoter regulates short-term memory (STM) retrieval in female mice only. **(A)** Mice were injected with either the nonfunctional control herpes simplex virus (HSV) (HSV-control) or HSV-*Cdk5*-zinc finger protein (ZFP)-p65 and subjected to fear conditioning (FC) followed by STM retrieval. **(B)** Male and **(C)** female FC mice injected with either HSV-*Cdk5*-ZFP-p65 or HSV-control showed gradual increase in freezing during acquisition (two-way analysis of variance, main effect of FC [$F_{4,24} = 110.8, p < .0001$], sex [$F_{3,18} = 1.303, p = .3042$], interaction [$F_{12,72} = 1.985, p = .0380$]). Male FC mice injected with HSV-*Cdk5*-ZFP-p65 did not differ in acquisition or short-term memory freezing from FC mice injected with HSV-control (preshock $p > .9999$, first shock $p > .9999$, second shock $p > .9999$, third shock $p > .9999$, and STM $p > .1188$). Female FC mice injected with HSV-*Cdk5*-ZFP-p65 displayed comparable freezing during acquisition and reduced freezing during STM retrieval (preshock $p > .9999$, first shock $p > .9999$, second shock $p > .9999$, third shock $p > .9999$, and STM $p = .0270$), compared with freezing of mice injected with HSV-control. Nonshocked, male control mice (C) injected with HSV-*Cdk5*-ZFP-p65 did not differ in freezing from control mice injected with HSV-control (preshock $p > .9999$, first shock $p > .9999$, second shock $p > .9999$, third shock $p > .9999$, and STM $p > .9999$). HSV-*Cdk5*-ZFP-p65-injected female mice froze less than male mice in STM retrieval ($p = .0068$). **(D, E)** Quantitative real-time polymerase chain reaction analysis of *Cdk5* after STM retrieval showed increased *Cdk5* expression in HSV-*Cdk5*-ZFP-p65-injected **(D)** male and **(E)** female mice (two-way analysis of variance, main effect of virus [$F_{1,7} = 6.795, p = .0351$], sex [$F_{1,7} = 0.3431, p = .5764$], interaction [$F_{1,7} = 2.263, p = .1762$]). *Cdk5* expression in mice injected with HSV-*Cdk5*-ZFP-p65 was greater than that of HSV-control-injected mice (male $p = .0011$, female $p = .0125$). *Cdk5* messenger RNA (mRNA) expression was increased in HSV-*Cdk5*-ZFP-p65-injected, nonshocked control mice after STM retrieval (two-way analysis of variance, main effect

To assess whether the reduction in LTM produced by HSV-*Cdk5*-ZFP-p65 in CA1 is specific to contextual fear memory, we analyzed the effect of this treatment on cocaine-induced CPP, another type of hippocampal-dependent learning and memory (49,50). Importantly, in the NAc of male mice, targeted methylation of *Cdk5* attenuates cocaine-induced CPP, while acetylation has no effect (15). We analyzed CPP behavior in male and female mice 7 days after CA1 injection of HSV-control and HSV-*Cdk5*-ZFP-p65 (Figure 6A). This time point corresponds to maximal acetylation during the preference test, which is analogous to maximal acetylation during the LTM retrieval in the fear conditioning paradigm. However, unlike the attenuated LTM retrieval observed in female mice following CA1 *Cdk5* promoter acetylation, we did not observe any effect of this manipulation on cocaine-induced CPP in mice of either sex (Figure 6B).

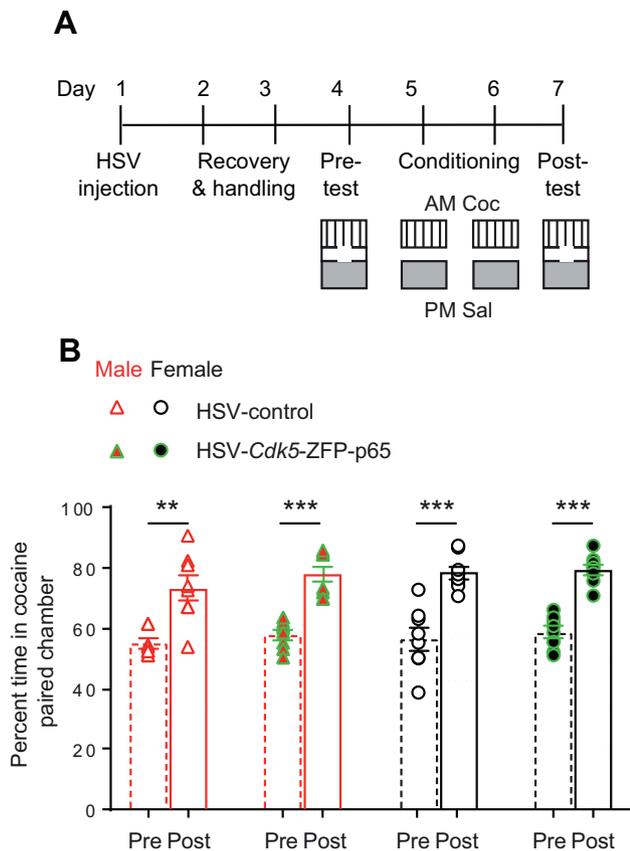


Figure 6. Epigenetic editing of cyclin-dependent kinase 5 (*Cdk5*) promoter in the hippocampus does not affect cocaine-induced conditioned place preference. **(A)** Timeline of herpes simplex virus (HSV) injection and cocaine-induced conditioned place preference. **(B)** Percent time spent in cocaine-paired chamber of virus-injected male and female mice. The male and female mice injected with either the nonfunctional control virus HSV-control or HSV-*Cdk5*-zinc finger protein (ZFP)-p65 spent more time in the cocaine-paired chamber (two-way analysis of variance, main effect of conditioning [$F_{1,7} = 53.17, p = .0002$], sex [$F_{3,21} = 2.518, p = .0858$]). The time spent in the cocaine-paired chamber was greater in all groups following conditioning (Bonferroni multiple comparison, male HSV-control $p = .0011$, female HSV-control $p = .0001$, male HSV-*Cdk5*-ZFP-p65 $p = .0005$, female HSV-*Cdk5*-ZFP-p65 $p = .0003$). All data are presented as mean \pm SEM. $n = 6-8$. $**p < .01$, $***p < .001$. AM Coc, morning cocaine; PM Sal, afternoon saline.

DISCUSSION

Sex differences in the extent and nature of mood disorders have led to a growing interest in the sexual specificity of molecular mechanisms underlying these diseases (51,52). Given our recent finding that histone acetylation and methylation of the *Cdk5* gene promoter regulate stress and reward responsiveness in male mice (15), we hypothesized that such mechanisms may be functionally relevant in female mice as well. We focused our attention on the role of *Cdk5* in fear-related memory in order to link the known functions of CDK5 in both learning- and stress-related behavior. Fear conditioning is a robust translational paradigm used in many cases to elucidate fear-related mechanisms of PTSD, such as fear extinction, fear inhibition, and generalization of fear (53-64). We applied the

innovative strategy of targeted epigenetic editing to elucidate the precise causal relevance of specific chromatin modifications to *Cdk5* expression and fear memory.

We first observed that female mice showed lower fear memory retrieval than did male mice, suggesting a female-specific mechanism for fear memory protection. While the difference is subtle, it is significant, reproducible, and consistent with the literature (65-68). We further found that enrichment of the activating hPTM, H3K9/14ac, at the *Cdk5* promoter did not change in female mice following fear conditioning, whereas in male mice, both H3K9/14ac and CDK5 expression were increased. To elucidate the causal relevance of male-specific *Cdk5* promoter acetylation, we targeted histone acetylation to the *Cdk5* promoter, which increased CDK5 expression in mice of both sexes. Surprisingly, *Cdk5* promoter acetylation attenuated STM and LTM retrieval in female mice only. This sexually dimorphic effect was accompanied by a female-specific increase in phosphorylation of tau protein, a CDK5 target implicated in learning and memory. Based on these findings, we propose a model in which *Cdk5* promoter acetylation, expression, and subsequent downstream target phosphorylation is sex-specifically regulated (Figure 7). We posit that in female mice, fear memory activation of *Cdk5* expression is blocked to control its downstream effects (e.g., tau protein phosphorylation) (Figure 7A). When this "break" on *Cdk5* expression is lifted by exogenous, locus-targeted acetylation, female mice display a fear memory deficit (Figure 7B). This finding is in contrast to what was found for male mice, in which both *Cdk5* promoter acetylation and activation are naturally increased following fear memory retrieval, and further, exogenous activation has no effect. Our model is supported by the fact that hyperphosphorylated tau protein affects microtubule dynamics, axonal transport, and neurite outgrowth, resulting in neurodegenerative pathologies (48). Female-specific neurological effects of hyperphosphorylated tau protein are well documented (44-48). For example, transgenic expression of hyperphosphorylated tau protein leads to a greater impairment in spatial learning and memory in female mice compared with that in male mice (45), and overexpression of corticotrophin-releasing factor increases tau protein phosphorylation in female mice, leading to an impairment in working memory (47). Given these sex differences, the "break" on fear-induced *Cdk5* activation in female CA1 may not be necessary in male CA1. While we investigated tau protein, *N*-methyl-D-aspartate receptor subunit 2B (NR2B) is another target of CDK5 associated with sex-specific synaptic transmission and plasticity (53,69). CDK5 phosphorylates NR2B and reduces its cell membrane expression (25). The literature reveals disagreement, however, on the precise mechanism of CDK5-mediated attenuation of fear memory. Recent studies point to the role of CDK5 in synaptic plasticity through both internalization (25) and reduced degradation of NR2B (70). Future studies to determine the extent of sex-specific downstream targeting by CDK5 will be useful to clarify the precise role of tau protein, NR2B, and other CDK5 targets in fear memory processing.

Sex differences in fear learning in rodents have been documented, with a role for calcium/calmodulin-dependent protein kinase alpha signaling in precipitating greater contextual fear conditioning in male rodents than in female rodents

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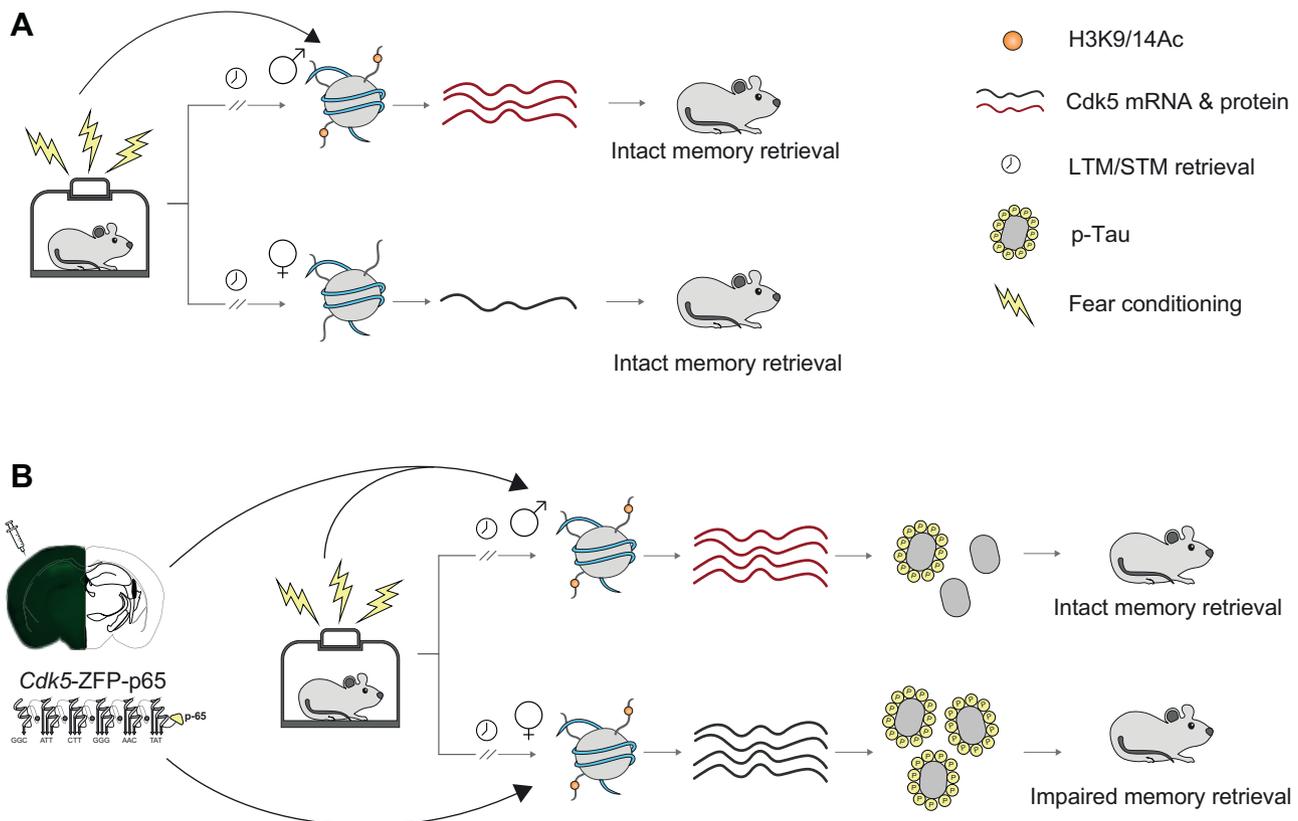


Figure 7. Our model proposes that cyclin-dependent kinase 5 (*Cdk5*) expression is sex-specifically regulated to control its downstream effects. **(A)** Fear conditioning of female mice does not change expression of CDK5 or enrichment of the activating histone posttranslational modification, histone H3 lysine 9/14 acetylation (H3K9/14ac), at the *Cdk5* promoter, whereas in male mice, fear conditioning activates CDK5 expression and enriches H3K9/14ac. **(B)** Locus-specific histone acetylation at the *Cdk5* promoter increases CDK5 expression in CA1 of both female and male mice. Yet this increase in CDK5 causes fear memory loss specifically in female mice, where it is correlated with a female-specific increase in phosphorylation of tau protein. We propose a model in which CDK5 activation in female CA1 is restrained following fear conditioning, to regulate the downstream effects of CDK5 activity. LTM, long-term memory; mRNA, messenger RNA; p-Tau, phosphorylated tau protein; STM, short-term memory; ZFP, zinc finger protein.

(71,72), which supports our similar observation. Additionally, heat exposure of female rats enhances fear extinction and retrieval associated with changes in hippocampal synaptic morphology (73). Sex differences have been observed in fear generalization, in which aversive experiences (e.g., shock) in one context cause unrelated neutral contexts to be processed as threatening. Male mice display greater c-Fos activity in the dorsal HPC during memory retrieval but less fear generalization, whereas female mice show greater fear generalization (74). One caveat to observed sex differences in fear-conditioning paradigms is the recent description of a “darting” phenotype—a rapid, forward movement across the chamber—associated with auditory fear conditioning in female, but not male, rats (75). Failure to quantify darting may contribute to the reduced retrieval measured in female mice. Beyond this, studies of sex-specific effects of fear conditioning have focused on the role of sex hormones (65), finding a role for estradiol in fear generalization (76) and for testosterone in auditory memory and long-term potentiation (77). We found no difference in *Cdk5* expression based on estrous cycle, but additional studies are needed to elucidate the interaction between basal sex hormone levels, estrous cycle stages, and fear memory retrieval in the epigenetic activation of CDK5 expression.

The lack of a fear memory deficit in male mice following *Cdk5* promoter acetylation was unexpected given that fear conditioning activated CDK5 expression and that conditional deletion of *Cdk5* in forebrain excitatory neurons results in poor spatial learning and memory in male mice (20). Such deletion is associated with hyperactivity, impaired cognitive function, and deficits in neurotransmitter release (20). Prior studies have used either pharmacological manipulation or conditional deletion of *Cdk5*, suggesting that the mechanism of manipulation may affect outcome measure. Targeted epigenetic editing recapitulated the endogenous mechanism and magnitude of CDK5 expression to reveal a previously unobserved, sex-specific epigenetic role for this kinase in fear memory retrieval. Importantly, we confirmed that CPP is not affected by *Cdk5* promoter acetylation in either CA1 or the NAc (15), indicating a specific role for CDK5 expression in contextual fear memory but not contextual reward memory. One conflicting outcome of these studies was the lack of CDK5 activation following fear conditioning of male mice injected with HSV-control. We reasoned that this deficit in *Cdk5* activation may be due to the effects of surgery and anesthesia in the context of viral manipulations but not in naïve mice. In support of this hypothesis, general anesthesia

is reported to cause histone modifications. For example, brains of 7-day-old rat pups show decreased histone acetylation following nitrous oxide and isoflurane anesthesia, while the suprachiasmatic nucleus of mice anesthetized with sevoflurane/oxygen shows decreased histone H4 acetylation of period circadian regulator 2 (37,78). We found that anesthesia reduced *Cdk5* expression, which could be overcome by HSV-*Cdk5*-ZFP-p65-activated expression. Fear conditioning alone (with HSV-control) was not sufficient to overcome repression of *Cdk5* expression by anesthesia and/or surgery, accounting for the lack of activation in this context.

The relevance of sex-specific gene expression to affective disorders is underscored in several studies of global and specific gene expression (23,79) and DNA methylation (79) in both male and female subjects. As reviewed recently (51), sex plays a key role in the extent of PTSD in the human population such that women develop this disorder at twice the rate of men (80). PTSD severity is associated with DNA methylation of the *SLC6A4* locus in female but not in male persons (6). PTSD may be associated with hormonally regulated DNA methylation of histone deacetylase 4, the expression of which is greater in the amygdala following auditory fear conditioning in wild-type female subjects compared with ovariectomized female subjects (8). In addition, increased promoter DNA methylation and reduced expression of the nuclear receptor subfamily 3 group C member 1 is associated with reduced PTSD risk in male, but not female, survivors of the Rwandan genocide (81). These compelling examples of sex-specific epigenetic regulation of DNA methylation at PTSD risk loci point to the potential for additional modes of gene regulation including hPTMs, like that identified in our study, as well as noncoding RNA (82–84) and chromatin conformation (85,86), mechanisms that are also implicated in susceptibility to stress disorders.

To contribute to understanding of sex-specific epigenetic regulation in fear memory, we specifically investigated the transcriptional regulation of CDK5 and found that the *Cdk5* promoter in male but not female CA1 is hyperacetylated after fear conditioning. Histone acetylation plays a vital role in gene expression related to fear memory (87–89) and anxiety (90–92). Sex-specific regulation of hPTMs has been documented, with male neonatal brain showing increased H3K9/14ac and H3K9me3 relative to female mice (93–95). Global chromatin profiling of female rat astrocytes finds a greater number of H3K4me3 peaks and greater H3K4-specific methyltransferase activity in young adult than in middle-aged female mice (96). There are also sex differences in the expression of histone acetyltransferases and deacetylases, which correlate with sex-specific gene expression (97). Quantification of histone acetyltransferase and/or histone deacetylase enrichment at *Cdk5*, as well as targeted deacetylation of *Cdk5*, can be applied to further our understanding of sex-specific epigenetic regulation of this locus.

Conclusions

Cdk5-targeted histone acetylation in CA1 attenuates fear memory retrieval in female, but not male, mice. This difference may be due to the increase in *Cdk5* expression and acetylation

in male, but not female, mice after fear memory retrieval, as well as female-specific tau protein phosphorylation following *Cdk5* activation. These results point to the relevance of *Cdk5* promoter acetylation in sexually dimorphic fear memory formation and related disorders.

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REFERENCES

- Day JJ, Sweatt JD (2012): Cognitive neuroepigenetics: A role for epigenetic mechanisms in learning and memory. *Neurobiol Learn Mem* 96:2–12.
- Borrelli E, Nestler EJ, Allis CD, Sassone-Corsi P (2008): Decoding the epigenetic language of neuronal plasticity. *Neuron* 96:1–974.
- Gräff J, Rei D, Guan J-S, Wang W-Y, Seo J, Hennig KM, et al. (2012): An epigenetic blockade of cognitive functions in the neurodegenerating brain. *Nature* 483:222–226.
- Lubin FD, Roth TL, Sweatt JD (2008): Epigenetic regulation of *BDNF* gene transcription in the consolidation of fear memory. *J Neurosci* 28:10576–10586.
- Malvaez M, Mhijaj E, Matheos DP, Palmery M, Wood MA (2011): CBP in the nucleus accumbens regulates cocaine-induced histone acetylation and is critical for cocaine-associated behaviors. *J Neurosci* 31:16941–16948.
- Nugent BM, Schwarz JM, McCarthy MM (2011): Hormones and behavior hormonally mediated epigenetic changes to steroid receptors in the developing brain: Implications for sexual differentiation. *Horm Behav* 59:338–344.
- Stevens JS, Almlil LM, Fani N, Gutman DA, Bradley B, Norrholm SD, et al. (2014): *PACAP* receptor gene polymorphism impacts fear responses in the amygdala and hippocampus. *Proc Natl Acad Sci U S A* 111:3158–3163.
- Maddox SA, Kilaru V, Shin J, Jovanovic T, Almlil LM, Dias BG, et al. (2018): Estrogen-dependent association of HDAC4 with fear in female mice and women with PTSD. *Mol Psychiatry* 23:658–665.
- Uddin M, Aiello AE, Wildman DE, Koenen KC, Pawelec G, de los Santos R, et al. (2010): Epigenetic and immune function profiles associated with posttraumatic stress disorder. *Proc Natl Acad Sci U S A* 107:9470–9475.
- Nestler EJ, Peña CJ, Kundakovic M, Mitchell A, Akbarian S (2016): Epigenetic basis of mental illness. *Neuroscientist* 22:447–463.
- Peter CJ, Akbarian S (2011): Balancing histone methylation activities in psychiatric disorders. *Trends Mol Med* 17:372–379.

Sexual Dimorphism of *Cdk5* Expression in Fear Memory

12. Geschwind DH (2009): Advances in autism. *Annu Rev Med* 60: 367–380.
13. Laplant Q, Vialou V, Covington HE, Dumitriu D, Feng J, Warren BL, *et al.* (2010): Dnmt3a regulates emotional behavior and spine plasticity in the nucleus accumbens. *Nat Neurosci* 13:1137–1143.
14. Heller EA, Cates HM, Peña CJ, Sun H, Shao N, Feng J, *et al.* (2014): Locus-specific epigenetic remodeling controls addiction- and depression-related behaviors. *Nat Neurosci* 17:1720–1727.
15. Heller EA, Hamilton PJ, Burek DD, Lombroso SI, Peña CJ, Neve RL, Nestler EJ (2016): Targeted epigenetic remodeling of the *Cdk5* gene in nucleus accumbens regulates cocaine- and stress-evoked behavior. *J Neurosci* 36:4690–4697.
16. Liu XS, Wu H, Krzisch M, Wu X, Graef J, Muffat J, *et al.* (2018): Rescue of Fragile X syndrome neurons by DNA methylation editing of the *FMR1* gene. *Cell* 172:979–991, e6.
17. Garriga-Canut M, Agustin-Pavon C, Herrmann F, Sanchez A, Diessen M, Fillat C, *et al.* (2012): Synthetic zinc finger repressors reduce mutant huntingtin expression in the brain of R6/2 mice. *Proc Natl Acad Sci U S A* 109:e3136–e3145.
18. Hawasli AH, Benavides DR, Nguyen C, Kansy JW, Hayashi K, Chambon P, *et al.* (2007): Cyclin-dependent kinase 5 governs learning and synaptic plasticity via control of NMDAR degradation. *Nat Neurosci* 10:880–886.
19. Zhong P, Liu X, Zhang Z, Hu Y, Liu SJ, Lezama-Ruiz M, *et al.* (2014): Cyclin-dependent kinase 5 in the ventral tegmental area regulates depression-related behaviors. *J Neurosci* 34:6352–6366.
20. Su SC, Rudenko A, Cho S, Tsai LH (2013): Forebrain-specific deletion of *Cdk5* in pyramidal neurons results in mania-like behavior and cognitive impairment. *Neurobiol Learn Mem* 105:54–62.
21. Rei D, Mason X, Seo J, Gräff J, Rudenko A, Wang J, *et al.* (2015): Basolateral amygdala bidirectionally modulates stress-induced hippocampal learning and memory deficits through a p25/*Cdk5*-dependent pathway. *Proc Natl Acad Sci U S A* 112:7291–7296.
22. Ramos-Miguel A, Meana JJ, Garcia-Sevilla JA (2013): Cyclin-dependent kinase-5 and p35/p25 activators in schizophrenia and major depression prefrontal cortex: Basal contents and effects of psychotropic medications. *Int J Neuropsychopharmacol* 16:683–689.
23. Labonté B, Engmann O, Purushothaman I, Menard C, Wang J, Tan C, *et al.* (2017): Sex-specific transcriptional signatures in human depression. *Nat Med* 23:1102–1111.
24. Bridi MS, Hawk JD, Chatterjee S, Safe S, Abel T (2017): Pharmacological activators of the NR4A nuclear receptors enhance LTP in a CREB/CBP-dependent manner. *Neuropsychopharmacology* 42:1243–1253.
25. Plattner F, Hernández A, Kistler TM, Pozo K, Zhong P, Yuen EY, *et al.* (2014): Memory enhancement by targeting *Cdk5* regulation of NR2B. *Neuron* 81:1070–1083.
26. Abel T, Nguyen PV, Barad M, Deuel TAS, Kandel ER, Bourchouladze R (1997): Genetic demonstration of a role for PKA in the late phase of LTP and in hippocampus-based long-term memory. *Cell* 88:615–626.
27. Duffy SN, Craddock KJ, Abel T, Nguyen PV (2001): Environmental enrichment modifies the PKA-dependence of hippocampal LTP and improves hippocampus-dependent memory. *Learn Mem* 8:26–41.
28. Vecsey CG, Hawk JD, Lattal KM, Stein JM, Fabian SA, Attner MA, *et al.* (2007): Histone deacetylase inhibitors enhance memory and synaptic plasticity via CREB: CBP-dependent transcriptional activation. *J Neurosci* 27:6128–6140.
29. Bibb JA, Chen J, Taylor JR, Svenningsson P, Nishi A, Snyder GL, *et al.* (2001): Effects of chronic exposure to cocaine are regulated by the neuronal protein *Cdk5*. *Nature* 410:376–380.
30. Calgioni CS (2009): Assessing reproductive status/stages in mice. *Curr Protoc Neurosci* 48:A41.1–A.41.8.
31. Perez SM, Chen L, Lodge DJ (2014): Alterations in dopamine system function across the estrous cycle of the MAM rodent model of schizophrenia. *Psychoneuroendocrinology* 47:88–97.
32. Kumar A, Choi KH, Renthal W, Tsankova NM, Theobald DEH, Truong HT, *et al.* (2005): Chromatin remodeling is a key mechanism underlying cocaine-induced plasticity in striatum. *Neuron* 48: 303–314.
33. Lu L, Grimm JW, Shaham Y, Hope BT (2003): Molecular neuro-adaptations in the accumbens and ventral tegmental area during the first 90 days of forced abstinence from cocaine self-administration in rats. *J Neurochem* 85:1604–1613.
34. Fischer A, Sananbenesi F, Schrick C, Spiess J, Radulovic J (2002): Cyclin-dependent kinase 5 is required for associative learning. *J Neurosci* 22:3700–3707.
35. Maze I, Covington HE, Dietz DM, LaPlant Q, Renthal W, Russo SJ, *et al.* (2010): Essential role of the histone methyltransferase G9a in cocaine-induced plasticity. *Science* 327:213–216.
36. Renthal W, Kumar A, Xiao G, Wilkinson M, Covington HE, Maze I, *et al.* (2009): Genome-wide analysis of chromatin regulation by cocaine reveals a role for sirtuins. *Neuron* 62:335–348.
37. Dalla Massara L, Osuru HP, Oklopčić A, Milanovic D, Joksimovic SM, Caputo V, *et al.* (2016): General anesthesia causes epigenetic histone modulation of c-Fos and brain-derived neurotrophic factor, target genes important for neuronal development in the immature rat hippocampus. *Anesthesiology* 124:1311–1327.
38. Ittnner LM, Ke YD, Delerue F, Bi M, Gladbach A, van Eersel J, *et al.* (2010): Dendritic function of tau mediates amyloid- β toxicity in alzheimer's disease mouse models. *Cell* 142:387–397.
39. Santacruz K, Lewis J, Spire T, Paulson J, Kotilinek L, Ingelsson M, *et al.* (2005): Medicine: Tau suppression in a neurodegenerative mouse model improves memory function. *Science* 309:476–481.
40. Lasagna-Reeves CA, Castillo-Carranza DL, Sengupta U, Guerrero-Munoz MJ, Kiritoshi T, Neugebauer V, *et al.* (2012): Alzheimer brain-derived tau oligomers propagate pathology from endogenous tau. *Sci Rep* 2:700.
41. Min SW, Chen X, Tracy TE, Li Y, Zhou Y, Wang C, *et al.* (2015): Critical role of acetylation in tau-mediated neurodegeneration and cognitive deficits. *Nat Med* 21:1154–1162.
42. Shentu YP, Huo Y, Feng XL, Gilbert J, Zhang Q, Liuyang ZY, *et al.* (2018): CIP2A causes tau/APP phosphorylation, synaptopathy, and memory deficits in Alzheimer's disease. *Cell Rep* 24:713–723.
43. Kimura T, Ishiguro K, Hisanaga S (2014): Physiological and pathological phosphorylation of tau by *Cdk5*. *Front Mol Neurosci* 7:65.
44. Köglberger S, Cordero-Maldonado ML, Antony P, Forster JI, Garcia P, Buttini M, *et al.* (2017): Gender-specific expression of ubiquitin-specific peptidase 9 modulates tau expression and phosphorylation: Possible implications for tauopathies. *Mol Neurobiol* 54:7979–7993.
45. Yue M, Hanna A, Wilson J, Roder H, Janus C (2011): Sex difference in pathology and memory decline in rTg4510 mouse model of tauopathy. *Neurobiol Aging* 32:590–603.
46. Devi L, Alldred MJ, Ginsberg SD, Ohno M (2010): Sex- and brain region-specific acceleration of β -amyloidogenesis following behavioral stress in a mouse model of Alzheimer's disease. *Mol Brain* 3:34.
47. Bangasser DA, Dong H, Carroll J, Plona Z, Ding H, Rodriguez L, *et al.* (2017): Corticotropin-releasing factor overexpression gives rise to sex differences in Alzheimer's disease-related signaling. *Mol Psychiatry* 22:1126–1133.
48. Cruz JC, Tseng HC, Goldman JA, Shih H, Tsai LH (2003): Aberrant *Cdk5* activation by p25 triggers pathological events leading to neurodegeneration and neurofibrillary tangles. *Neuron* 30:471–483.
49. Sjulson L, Peyrache A, Cumpelik A, Cassataro D, Buzsáki G (2018): Cocaine place conditioning strengthens location-specific hippocampal coupling to the nucleus accumbens. *Neuron* 98:926–934, e5.
50. Meyers RA, Zavala AR, Speer CM, Neisewander JL (2006): Dorsal hippocampus inhibition disrupts acquisition and expression, but not consolidation, of cocaine conditioned place preference. *Behav Neurosci* 120:401–412.
51. Ramikie TS, Ressler KJ (2018): Mechanisms of sex differences in fear and posttraumatic stress disorder. *Biol Psychiatry* 83:876–885.
52. McCarthy MM, Nugent BM, Lenz KM (2017): Neuroimmunology and neuroepigenetics in the establishment of sex differences in the brain. *Nat Rev Neurosci* 18:471–484.
53. Zeng Y, Lv X, Zeng S, Shi J (2009): Activity-dependent neuronal control of gap-junctional communication in fibroblasts. *Brain Res* 1280:13–22.

54. Gourley SL, Kedves AT, Olausson P, Taylor JR (2009): A history of corticosterone exposure regulates fear extinction and cortical NR2B, GluR2/3, and BDNF. *Neuropsychopharmacology* 34:707–716.
55. Mahan AL, Ressler KJ (2013): Fear conditioning, synaptic plasticity, and the amygdala: Implications for posttraumatic stress disorder. *Trends Neurosci* 35:24–35.
56. Ding AY, Li Q, Zhou IY, Ma SJ, Tong G, McAlonan GM, Wu EX (2013): MR diffusion tensor imaging detects rapid microstructural changes in amygdala and hippocampus following fear conditioning in mice. *PLoS One* 8:e51704.
57. Briscione MA, Jovanovic T, Norrholm SD (2014): Conditioned fear associated phenotypes as robust, translational indices of trauma-, stressor-, and anxiety-related behaviors. *Front Psychiatry* 5:88.
58. Johansen JP, Cain CK, Ostroff LE, Ledoux JE (2011): Molecular mechanisms of fear learning and memory [published correction appears in *Cell* 2011;147:948]. *Cell* 147:509–524.
59. Amstadter AB, Nugent NR, Koenen KC (2009): Genetics of PTSD: Fear conditioning as a model for future research. *Psychiatr Ann* 39:358–367.
60. Chang C, Knapska E, Orsini CA, Rabinak CA, Zimmerman JM, Maren S (2009): Fear extinction in rodents. *Curr Protoc Neurosci* 47:8.23.1–8.23.17.
61. Lissek S, Powers AS, McClure EB, Phelps EA, Woldehawariat G, Grillon C, Pine DS (2005): Classical fear conditioning in the anxiety disorders: A meta-analysis. *Behav Res Ther* 43:1391–1424.
62. Rubin DC, Berntsen D, Bohni MK (2008): A memory-based model of posttraumatic stress disorder: evaluating basic assumptions underlying the PTSD diagnosis. *Psychol Rev* 115:985–1011.
63. Morey RA, Dunsmoor JE, Haswell CC, Brown VM, Vora A, Weiner J, et al. (2015): Fear learning circuitry is biased toward generalization of fear associations in posttraumatic stress disorder. *Transl Psychiatry* 5:e700.
64. Kaouane N, Porte Y, Vallée M, Brayda-Bruno L, Mons N, Calandreau L, et al. (2012): Glucocorticoids can induce PTSD-like memory impairments in mice. *Science* 335:1510–1513.
65. Lynch J, Cullen PK, Jasnow AM, Riccio DC (2013): Sex differences in the generalization of fear as a function of retention intervals. *Learn Mem* 20:628–632.
66. Inslicht SS, Metzler TJ, Garcia NM, Pineles SL, Milad MR, Orr SP, et al. (2013): Sex differences in fear conditioning in posttraumatic stress disorder. *J Psychiatr Res* 47:64–71.
67. Lebron-Milad K, Abbs B, Milad MR, Linnman C, Rougemont-Bücking A, Zeidan MA, et al. (2012): Sex differences in the neurobiology of fear conditioning and extinction: A preliminary fMRI study of shared sex differences with stress-arousal circuitry. *Biol Mood Anxiety Disord* 2:7.
68. Koss WA, Frick KM (2017): Sex differences in hippocampal function. *J Neurosci Res* 95:539–562.
69. Boon WC, Diepstraten J, Van Der Burg J, Jones MEE, Simpson ER, Van Den Buuse M (2005): Hippocampal NMDA receptor subunit expression and watermaze learning in estrogen deficient female mice. *Brain Res Mol Brain Res* 140:127–132.
70. Hawasli AH, Bibb JA (2007): Alternative roles for Cdk5 in learning and synaptic plasticity. *Biotechnol J* 2:941–948.
71. Mizuno K, Ris L, Sanchez-Capelo A, Godaux E, Giese KP (2006): Ca²⁺/calmodulin kinase kinase is dispensable for brain development but is required for distinct memories in male, though not in female, mice. *Mol Cell Biol* 26:9094–9104.
72. Mizuno K, Dempster E, Mill J, Giese KP (2012): Long-lasting regulation of hippocampal *Bdnf* gene transcription after contextual fear conditioning. *Genes Brain Behav* 11:651–659.
73. Gruene TM, Lipps J, Rey CD, Bouck A, Shansky RM (2014): Neurobiology of learning and memory heat exposure in female rats elicits abnormal fear expression and cellular changes in prefrontal cortex and hippocampus. *Neurobiol Learn Mem* 115:38–42.
74. Keiser AA, Turnbull LM, Darian MA, Feldman DE, Song I, Tronson NC (2017): Sex differences in context fear generalization and recruitment of hippocampus and amygdala during retrieval. *Neuropsychopharmacology* 42:397–407.
75. Gruene TM, Flick K, Stefano A, Shea SD, Shansky RM (2015): Sexually divergent expression of active and passive conditioned fear responses in rats. *Elife* 4:e11352.
76. Lynch JF, Dejanovic D, Winiacki P, Mulvany J, Ortiz S, Riccio DC, Jasnow AM (2014): Activation of ER β modulates fear generalization through an effect on memory retrieval. *Horm Behav* 66:421–429.
77. Chen LS, Tzeng WY, Chuang JY, Cherng CG, Gean PW, Yu L (2014): Roles of testosterone and amygdaloid LTP induction in determining sex differences in fear memory magnitude. *Horm Behav* 66:498–508.
78. Mori K, Iijima N, Higo S, Aikawa S, Matsuo I, Takumi K, et al. (2014): Epigenetic suppression of mouse *Per2* expression in the suprachiasmatic nucleus by the inhalational anesthetic, sevoflurane. *PLoS One* 9:E87319.
79. Nugent BM, Wright CL, Shetty AC, Hodes GE, Lenz KM, Mahurkar A, et al. (2015): Brain feminization requires active repression of masculinization via DNA methylation. *Nat Neurosci* 18:690–697.
80. Haskell SG, Gordon KS, Mattocks K, Duggal M, Erdos J, Justice A, Brandt CA (1988): Psychological unemployment: A selected overview. *J R Soc Health* 108:29–31, 33.
81. Vukojevic V, Kolassa I, Fastenrath M, Gschwind L, Spalek K, Milnik A, et al. (2014): Epigenetic modification of the glucocorticoid receptor gene is linked to traumatic memory and post-traumatic stress disorder risk in genocide survivors. *J Neurosci* 34:10274–10284.
82. Morgan CP, Bale TL (2011): Early Prenatal Stress Epigenetically Programs Dysmasculinization in Second-Generation Offspring via the Paternal Lineage. *J Neurosci* 31:11748–11755.
83. Rodgers AB, Morgan CP, Bronson SL, Revello S, Bale TL (2013): Paternal stress exposure alters sperm microRNA content and reprograms offspring HPA stress axis regulation. *J Neurosci* 33:9003–9012.
84. Hodes GE, Pfau ML, Purushothaman I, Ahn HF, Golden SA, Christoffel DJ, et al. (2015): Sex differences in nucleus accumbens transcriptome profiles associated with susceptibility versus resilience to subchronic variable stress. *J Neurosci* 35:16362–16376.
85. Klengel T, Mehta D, Anacker C, Rex-Haffner M, Pruessner JC, Pariante CM, et al. (2013): Allele-specific FKBP5 DNA demethylation mediates gene-childhood trauma interactions. *Nat Neurosci* 16:33–41.
86. Mitchell AC, Javidfar B, Bicks LK, Neve R, Garbett K, Lander SS, et al. (2016): Longitudinal assessment of neuronal 3D genomes in mouse prefrontal cortex. *Nat Commun* 7:12743.
87. Peixoto L, Abel T (2013): The role of histone acetylation in memory formation and cognitive impairments. *Neuropsychopharmacology* 38:62–76.
88. Mews P, Donahue G, Drake AM, Luczak V, Abel T, Berger SL (2017): Acetyl-CoA synthetase regulates histone acetylation and hippocampal memory. *Nature* 546:381–386.
89. Levenson JM, O'Riordan KJ, Brown KD, Trinh MA, Molfese DL, Sweatt JD (2004): Regulation of histone acetylation during memory formation in the hippocampus. *J Biol Chem* 279:40545–40559.
90. Weaver ICG, Cervoni N, Champagne FA, D'Alessio AC, Sharma S, Seckl JR, et al. (2004): Epigenetic programming by maternal behavior. *Nat Neurosci* 7:847–854.
91. Hollis F, Duclot F, Gunjan A, Kabbaj M (2011): Individual differences in the effect of social defeat on anhedonia and histone acetylation in the rat hippocampus. *Horm Behav* 59.
92. Gundersen BB, Blendy JA (2009): Effects of the histone deacetylase inhibitor sodium butyrate in models of depression and anxiety. *Neuropharmacology* 57:67–74.
93. Bhadra U, Pal-Bhadra M, Birchler JA (2000): Histone acetylation and gene expression analysis of sex lethal mutants in drosophila. *Genetics* 155:753–763.
94. Champagne FA, Weaver ICG, Diorio J, Dymov S, Szyf M, Meaney MJ (2006): Maternal care associated with methylation of the estrogen receptor- α 1b promoter and estrogen receptor- α expression in the medial preoptic area of female offspring. *Endocrinology* 147:2909–2915.
95. Tsai H-W, Grant P a, Rissman EF (2009): Sex differences in histone modifications in the neonatal mouse brain. *Epigenetics* 4:47–53.
96. Chisholm NC, Henderson ML, Selvamani A, Park MJ, Dindot S, Miranda RC, Sohrabji F (2015): Histone methylation patterns in astrocytes are influenced by age following ischemia. *Epigenetics* 10:142–152.
97. Murray EK, Hien A, De Vries GJ, Forger NG (2009): Epigenetic control of sexual differentiation of the bed nucleus of the stria terminalis. *Endocrinology* 150:4241–4247.