

# Hippocampal HDAC4 Contributes to Postnatal Fluoxetine-Evoked Depression-Like Behavior

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Fluoxetine treatment in adulthood evokes antidepressant and anxiolytic responses. Paradoxically, postnatal fluoxetine (PNFlx) induces persistent depression- and anxiety-like behaviors. The mechanistic underpinnings of this paradox remain poorly understood. Here, we examined specific molecular changes in the rat hippocampus that accompany perturbed emotionality observed across life following PNFlx. PNFlx-induced hippocampal gene regulation observed in microarray and quantitative PCR studies indicate functional enrichment of genes involved in response to organic substances, protein kinase pathways, DNA binding, and transcriptional repression. We noted specific transcripts (*Hdac4*, mammalian target of rapamycin (*mTOR*), *Gnail*, protein kinase C gamma (*Prkcc*), and hyperpolarization-activated cyclic nucleotide-gated channel I (*Hcn1*)) that were consistently dysregulated across life, and selectively influenced by postnatal, but not adult, fluoxetine. Increased histone deacetylase-4 (HDAC4) recruitment, accompanied by decreased activating histone acetylation marks at the *mTOR* and *Gnail* promoters, indicate a role for HDAC4 in PNFlx-mediated gene dysregulation. Strikingly, coadministration of the HDAC inhibitor sodium butyrate with PNFlx prevented the dysregulation of *Hdac4* and *mTOR*, and the emergence of depression- and anxiety-like behavior. Importantly, we also find that retreatment of PNFlx animals with fluoxetine in adulthood reversed the increased *Hdac4* expression, prevented HDAC4 recruitment to the *mTOR* and *Gnail* promoters, and attenuated the decline in *mTOR* and *Gnail* expression, coincident with normalization of PNFlx-evoked depression- and anxiety-like behavior. Further, we show that viral-mediated hippocampal overexpression of *Hdac4* was sufficient to induce depression-, but not anxiety-, like behavior in adulthood. Our results highlight the unique nature of molecular signatures evoked by PNFlx, and implicate HDAC4 in the dysregulated gene expression and emergence of perturbed emotionality following fluoxetine exposure in early life.

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## INTRODUCTION

Fluoxetine (Prozac), a selective serotonin reuptake inhibitor (SSRI), is the drug of choice for gestational/postpartum depression (Ward and Zamorski, 2002) and children/adolescents with mood disorders (Kodish *et al*, 2011), given its perceived favorable risk to benefit ratio. However, preclinical and clinical studies have raised concerns about early fluoxetine exposure (Ansorge *et al*, 2004; Olivier *et al*, 2011). Studies in mouse models indicate enhanced depression- and anxiety-like behavior following postnatal SSRI administration (Ansorge *et al*, 2004, 2008; Karpova *et al*, 2009; Popa *et al*, 2008). Perinatal SSRI administration disrupts sleep (Popa *et al*, 2008), circadian rhythms (Kiryanova *et al*, 2013), stress responses (Norcross *et al*, 2008), and serotonin neurocircuitry (Silva *et al*, 2010;

Maciag *et al*, 2006). Similarly, deletion of the serotonin transporter, the molecular target of SSRIs, early in life induces depression- and anxiety-like symptoms in mice (Holmes *et al*, 2003; Lira *et al*, 2003). Clinical evidence supports an association between pediatric and adolescent fluoxetine usage and adverse effects on emotionality (Olivier *et al*, 2011). In contrast, adult fluoxetine (AFlx) induces antidepressant and anxiolytic responses in patients (Brambilla *et al*, 2005; Olivier *et al*, 2011) and animal models (Dulawa *et al*, 2004). These studies suggest opposing effects of fluoxetine on emotionality based on the temporal window of administration, for which the underlying mechanisms are unknown.

We hypothesized that depression- and anxiety-like behaviors in animals with a history of postnatal fluoxetine (PNFlx) involve underlying molecular changes maintained long after drug treatment. We examined PNFlx-induced molecular changes in the hippocampus, a limbic structure implicated in the regulation of mood (Deacon *et al*, 2002; Kheirbek *et al*, 2013) and known to transiently express the serotonin transporter during postnatal life (Narboux-Nême *et al*, 2008). We identified unique molecular signatures associated with postnatal, but not adult, fluoxetine treat-

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ment, including a persistent upregulation of the histone-modifying enzyme, *Hdac4* (histone deacetylase-4). PNFlx-evoked depression- and anxiety-like behavior, and specific molecular signatures were prevented by HDAC inhibitor co-treatment during postnatal life, and also normalized by fluoxetine retreatment in adulthood. Further, hippocampal HDAC4 overexpression was sufficient to induce depression-like behavior. Our findings implicate hippocampal HDAC4 as a mediator of the persistent increases in depression-like behavior following PNFlx.

## MATERIALS AND METHODS

### Animals

Male Sprague–Dawley rats bred at the Tata Institute of Fundamental Research (TIFR) animal facility were used for experiments and maintained with *ad libitum* access to food and water. Procedures followed the National Institutes of Health Guide for the Care and Use of Animals, and were approved by the TIFR Animal Ethics Committee. Viral overexpression experiments were performed at the Mount Sinai School of Medicine and were approved by the Institutional Animal Care and Use Committee.

### Drug Treatments

PNFlx treatment involved oral fluoxetine (10 mg/kg, Sigma, USA) or vehicle (5% sucrose) administration to rat pups from postnatal day (P) 2 to P21. AFlx treatment involved once daily intraperitoneal administration of fluoxetine (10 mg/kg) or vehicle (saline) for 21 days to 2-month-old animals. To address whether AFlx reversed PNFlx-evoked changes there were three groups: control (Ctrl), PNFlx, and PNFlx + AFlx. Behavioral experiments commenced after 21 days of AFlx and fluoxetine treatment was continued through behavioral experiments. For HDAC inhibitor sodium butyrate (SB) experiments, pups received fluoxetine (10 mg/kg), fluoxetine (10 mg/kg) + SB (300 mg/kg, Sigma), or vehicle (5% sucrose) orally from P2 to 21 with three groups: Ctrl, PNFlx, and PNFlx + PNSB. Drug doses were based on prior literature (Dulawa *et al*, 2004; Ansorge *et al*, 2008; Kim *et al*, 2009).

### Behavioral Tests

Behavioral tests were performed at specific time points across life. Pair-wise social play was assessed in Ctrl and PNFlx rats (P28–P32;  $n = 8–10$  per group) by an investigator blind to treatment conditions (Homberg *et al*, 2007). Depression- and anxiety-like behavior in Ctrl and PNFlx groups (2 months;  $n = 8–10$  per group) were assessed using the modified forced swim test (FST), open field test (OFT), and elevated plus maze (EPM). Further, Ctrl and PNFlx animals at 18 months of age ( $n = 8–10$  per group), postnatal SB experiment ( $n = 6–12$  per group), AFlx retreatment experiment ( $n = 6–10$  per group), and hippocampal HDAC4 overexpression experiment ( $n = 6–7$  per group) were tested on the OFT and FST. For detailed methods and behavioral paradigms (Supplementary Figure S1), see Supplementary Information. Behaviors were analyzed using Ethovision 3.1 (Noldus, the Netherlands).

### Microarray Experiments

Microarray analysis with a custom rat array  $8 \times 15$  K was performed on hippocampi from control and PNFlx animals (2 months,  $n = 4$  per group) as previously described (Benekareddy *et al*, 2010). Data were extracted using Agilent Feature Extraction software v 10.5.1.1 and differentially regulated genes were filtered ( $> 1.35$  for upregulation and  $< 0.74$  for downregulation). Statistical analysis was performed using *t*-test ( $p < 0.05$ ) with multiple comparison corrections (Benjamini and Hochberg, 1995). Array data are deposited in the NCBI Gene Expression Omnibus (Accession number GSE42940). Functional analysis was performed using DAVID (Database for Annotation Visualization and Integrated Discovery) on array data after removal of odorant receptors as potential false positives to avoid any bias to gene ontology analysis (Dennis *et al*, 2003).

### Chromatin Immunoprecipitation Assay

ChIP analysis was performed as previously described (Renthal *et al*, 2007). ChIP assays measured H3 (H3ac) and H4 (H4ac) acetylation, and HDAC4 enrichment at gene promoters. Chromatin enriched for H3ac, H4ac, and HDAC4 was subjected to quantitative PCR (qPCR) analysis (see Supplementary Information).

### Quantitative PCR

qPCR was performed on Ctrl and PNFlx (P21, 2 months and 18 months;  $n = 8–10$  per group); Ctrl and AFlx ( $n = 6–10$  per group); Ctrl, PNFlx, and PNFlx + AFlx ( $n = 6–11$  per group); Ctrl, PNFlx, and PNFlx + PNSB ( $n = 6–10$  per group); HSV-GFP and HSV-HDAC4-GFP ( $n = 4$  per group); and qPCR for ChIP analysis in Ctrl, PNFlx, and PNFlx + AFlx ( $n = 6–10$  per group). Reverse transcribed RNA was subjected to qPCR using primers for genes of interest (Supplementary Figures S2 and S3). Quantification was done using the  $\Delta\Delta C_t$  method as previously described (Bookout and Mangelsdorf, 2003) and data were normalized to hypoxanthine-guanine phosphoribosyltransferase (*Hprt1*), which was not regulated by any of the treatments used in this study. Results were compared with the controls and expressed as fold change  $\pm$  SEM.

### Western Blot Analysis

Hippocampi were lysed and separated by SDS–PAGE, and transferred to polyvinylidene difluoride membranes (GE Healthcare, Buckinghamshire, UK). Membranes were blocked, incubated with rabbit anti-mTOR (1:1000, Cell Signaling Technologies), rabbit anti-HDAC4 (1:500, Abcam, UK), rabbit anti-GAPDH (1:1000, Cell Signaling Technologies), and rabbit anti- $\beta$ III-tubulin antibody (1:1000; Covance, Princeton, NJ) before incubation with a horseradish peroxidase-conjugated goat anti-rabbit secondary antibody (GE Healthcare) and detection with an ECL substrate (GE Healthcare). Densitometric measurements were performed using a Macintosh-based Scion Image software (Scion, USA) and levels of mTOR and HDAC4 were normalized to  $\beta$ III-tubulin and GAPDH, respectively.

## Viral-Mediated Gene Transfer

A bicistronic HSV-HDAC4-GFP and HSV-GFP (Clark *et al*, 2002; Renthal *et al*, 2007) virus with expression driven off an IE4/5 promoter (HDAC4 cDNA was a kind gift of E Olson, University of Texas Southwestern Medical Center) were bilaterally delivered into the hippocampus of treatment-naïve adult rats as previously described (see Supplementary Information). Viral targeting and overexpression were assessed by imaging GFP fluorescence, qPCR, and western blot analysis.

## Statistical Analysis

Experiments with two or three groups were analyzed using unpaired Student's *t*-test or one-way ANOVA followed by Student–Newman–Keul's *post hoc* test, respectively, with significance at  $p < 0.05$  (InStat, Graphpad Software, USA).

## RESULTS

### PNFlx Evokes Persistent Depression- and Anxiety-Like Behavior

We examined the consequences of PNFlx treatment on juvenile play (P28–32) and depression- and anxiety-like behavior in adulthood and at 18 months of age (Figure 1a, Supplementary Figure S1). Adolescent PNFlx animals showed decreased social grooming and frequency of pouncing (Figure 1b and c), whereas follow and chase was unaltered (Figure 1d). PNFlx animals in adulthood showed depression- and anxiety-like behavior on the FST (Figure 1e–g), OFT (Figure 1h–j), and EPM (Supplementary Figure S5). We noted increased immobility time, decreased latency to the first immobility event, and a trend ( $p = 0.07$ ) toward increased immobility events on the FST in PNFlx animals. On the OFT, PNFlx animals exhibited an increased latency to approach the center, decreased percent time and path length in the center of the arena. Total distance traversed in the OFT was unaltered (Supplementary Figure S4). Depression- and anxiety-like behavior on the FST (Figure 1k and l, Supplementary Figure S6) and OFT (Figure 1m, Supplementary Figure S6) persisted up to 18 months in PNFlx animals. We observed no change in hippocampal-dependent spatial learning in PNFlx animals in adulthood (Supplementary Figure S7). Our results indicate that mood-related behavioral changes emerge as early as adolescence in PNFlx animals, and persist up to 18 months of age.

### PNFlx Alters the Hippocampal Transcriptome

Microarray analysis on hippocampi from 2-month-old PNFlx animals (Figure 1n) revealed significant enrichment for genes associated with response to organic substances and glucocorticoid stimuli, regulation of cellular protein metabolic processes and protein kinase activity, DNA binding, and transcription repressor activity (Figure 1o, Supplementary Figure S8). We noted a preponderance of downregulated genes (771), as compared with those exhibiting a significant upregulation (519). qPCR validation in an independent cohort of 2-month-old PNFlx animals (Figure 2a) indicated strong reproducibility of our micro-

array results (Figure 2c). qPCR analysis confirmed the regulation of multiple array-identified targets, including *Hdac4*, *mTOR*, *Prkcc*, *Gnai1*, and *Hcn1*.

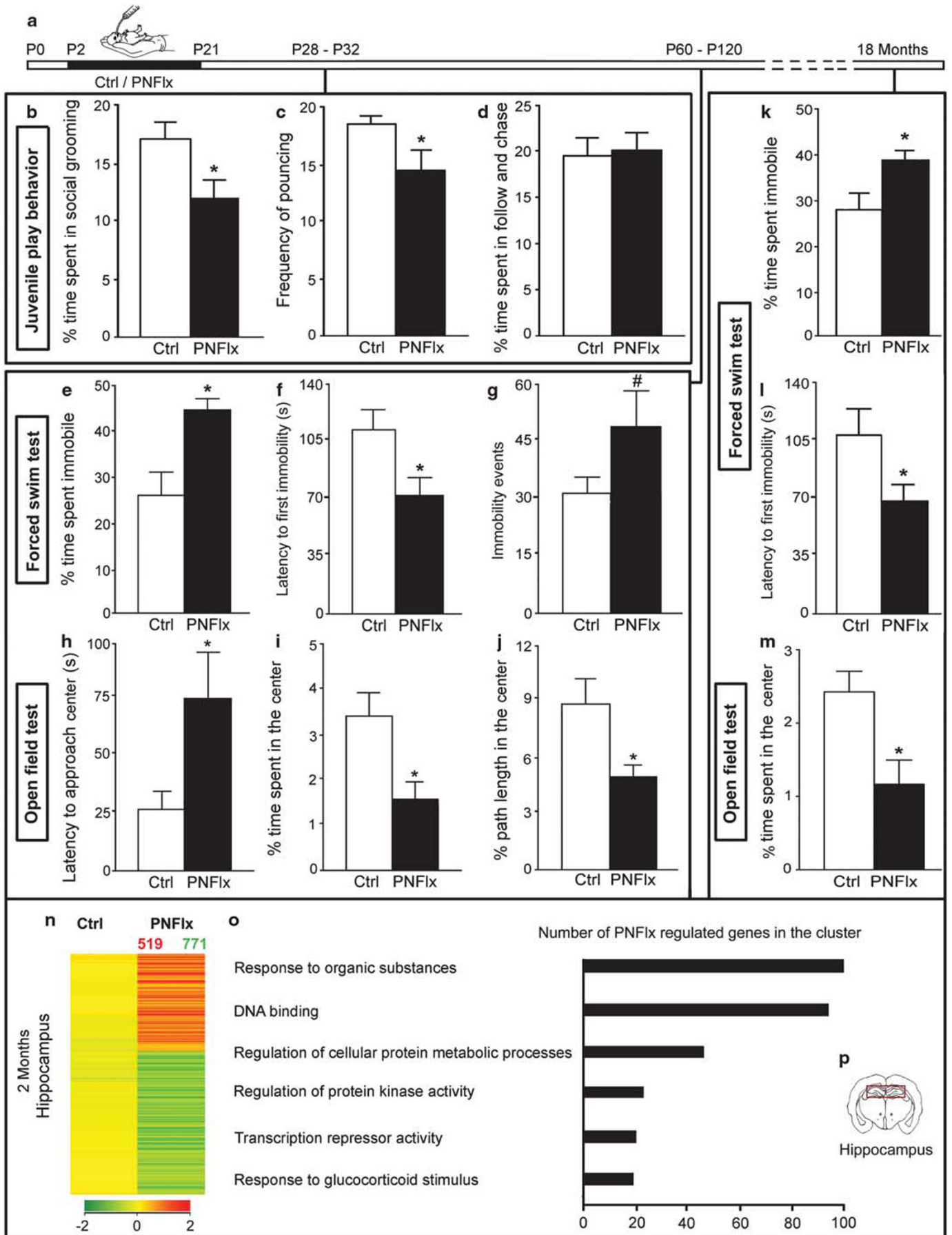
### PNFlx Induces Dysregulation of Specific Genes in the Hippocampus Across Life

We next examined whether the PNFlx-evoked gene dysregulation in adulthood was observed across life, in keeping with the persistent alterations in behavior. Strikingly, *Hdac4* expression was significantly upregulated at all three ages examined (P21, 2 months and 18 months) in PNFlx animals (Figure 2a–d). Although other upregulated genes validated in 2-month-old PNFlx animals did not exhibit a stable pattern, specific downregulated genes (*mTOR*, *Gnai1*, *Prkcc*, and *Hcn1*) showed a persistent decline in expression across life (Figure 2b–d). To address whether *Hdac4*, *mTOR*, *Gnai1*, *Prkcc*, and *Hcn1* gene regulation was unique to PNFlx treatment, which evokes depression- and anxiety-like behavior, we examined the influence of AFlx known to induce antidepressant and anxiolytic responses on the expression of these genes. AFlx treatment did not alter hippocampal *Hdac4*, *mTOR*, *Gnai1*, *Prkcc*, or *Hcn1* expression (Figure 2e and f). Conversely, PNFlx did not influence hippocampal *Bdnf*, *TrkB*, *Gdnf*, *c-Fos*, or *Arc* expression, which are regulated following AFlx (Nibuya *et al*, 1995; Hisaoka *et al*, 2001; Miller *et al*, 2008; Figure 2g). The effects of PNFlx were selective to *Hdac4* with no change observed in other *Hdacs* examined (Figure 2h). The enhanced *Hdac4* expression in PNFlx animals in adulthood was accompanied by a significant enrichment of H3ac and H4ac at the *Hdac4* promoter (Figure 2i). These results implicate persistent dysregulation of *Hdac4*, *mTOR*, *Gnai1*, *Prkcc*, and *Hcn1* as molecular signatures that accompany the life-long behavioral changes evoked by PNFlx.

As we observed a persistent decline in *mTOR*, *Gnai1*, *Prkcc*, and *Hcn1* expression in PNFlx animals, we asked whether HDAC4, a potent transcriptional repressor, contributes to the decline in expression of these genes. ChIP analysis indicated significant HDAC4 enrichment at *mTOR* and *Gnai1* (Figure 2j), but not *Prkcc* and *Hcn1* (Supplementary Figure S9), promoters following PNFlx. HDAC4 enrichment was accompanied by a significant decline in H4ac at *mTOR* and *Gnai1* (Figure 2k), but not H3ac at the *mTOR* and *Gnai1* promoters (Supplementary Figure S10).

### Postnatal Treatment with SB Prevents the Emergence of PNFlx-Evoked Molecular and Behavioral Changes

To test the hypothesis that HDAC4 may contribute to PNFlx-evoked behavioral outcomes, animals were coadministered the HDAC inhibitor, SB, with fluoxetine in postnatal life. We profiled depression- and anxiety-like behavior and gene expression changes in adulthood (Figure 3a). Combined PNFlx and SB treatment (PNFlx + PNSB) prevented the emergence of depression- and anxiety-like behavior on the FST (Figure 3b–e) and OFT (Figure 3f–j). One-way ANOVA and *post hoc* analysis indicated that the PNFlx + PNSB group differed significantly from PNFlx animals on immobility time (Figure 3c) and latency to first immobility (Figure 3d), with no differences on immobility events (Figure 3e). On the OFT, the PNFlx + PNSB group did not



show decreases in percent time, path length, and number of visits to the center of the arena observed in PNFlx animals, and were similar to controls (Figure 3f–j). A history of PNSB treatment itself did not alter behavior on the OFT or FST tasks (Supplementary Figure S11) in adulthood.

We examined whether SB treatment could similarly prevent the PNFlx-mediated dysregulation of specific genes within the hippocampus. PNFlx + PNSB animals did not exhibit *Hdac4* upregulation (Figure 3k) and failed to show the PNFlx-associated *mTOR* decline (Figure 3l). In contrast, decreased *Gnai1*, *Prkcc*, and *Hcn1* expression (Figure 3m–o) was still observed in the PNFlx + PNSB group. These results indicate that concomitant HDAC inhibition prevents the emergence of PNFlx-evoked depression- and anxiety-like behaviors, as well as the regulation of specific genes.

### AFlx Treatment Reverses Behavioral and Molecular Signatures Associated with PNFlx

Given that PNFlx and AFlx evoke distinct molecular and behavioral changes, we examined whether retreatment with fluoxetine in adulthood could influence PNFlx-evoked outcomes (Figure 4a). One-way ANOVA analysis indicated that AFlx completely reversed the PNFlx-evoked depression- and anxiety-like behaviors in the FST and OFT (Figure 4b–e). Further, PNFlx + AFlx animals exhibited antidepressant-like FST behaviors with decreased percent immobility and immobility events (Figure 4b and c). Enhanced histone acetylation at the *Hdac4* promoter and increased hippocampal *Hdac4* mRNA were reversed to control levels in PNFlx + AFlx animals (Figure 4f and g). PNFlx + AFlx animals did not exhibit increased HDAC4 recruitment at the *Gnai1* and *mTOR* promoters (Figure 4h and j), and showed attenuation of the PNFlx-evoked decline in *Gnai1* and *mTOR* (Figure 4i and k) expression. Western blot analysis revealed decreased mTOR levels in hippocampi from PNFlx animals, which was ameliorated in the PNFlx + AFlx group (Figure 4l). These results indicate that retreatment with fluoxetine in adulthood rescues behavioral and molecular changes evoked by PNFlx.

### Hippocampal *Hdac4* Overexpression in Adulthood Evokes Depression-Like Behavior

We next examined whether viral HDAC4 overexpression in adulthood could recapitulate PNFlx-evoked behavioral phenotypes. We performed bilateral stereotaxic delivery of HSV-HDAC4-GFP or HSV-GFP into hippocampi of treatment-naïve adult animals and assessed OFT and FST behavior (Figure 5a). HSV-HDAC4-GFP localization was

confirmed by GFP fluorescence (Figure 5b and c). HDAC4 overexpression was indicated by a strong trend toward significance in both qPCR ( $p=0.07$ ) and western blot ( $p=0.062$ ) analysis (Figure 5d and e). Hippocampal HDAC4 overexpression evoked a significant decline in *Gnai1* expression and a strong trend ( $p=0.051$ ) toward a decline in *mTOR* expression (Figure 5d).

HSV-HDAC4-GFP animals exhibited enhanced percent immobility, decreased latency to first immobility, and a strong trend ( $p=0.07$ ) for increased immobility events (Figure 5f–h). On the OFT, the HSV-HDAC4-GFP group did not differ from controls (percent time: HSV-GFP =  $5.27 \pm 0.86$ , HSV-HDAC4-GFP =  $9.49 \pm 2.49$ ; percent path length: HSV-GFP =  $5.76 \pm 0.79$ , HSV-HDAC4-GFP =  $7.93 \pm 3.10$ ; number of visits: HSV-GFP =  $28.8 \pm 4.7$ , HSV-HDAC4-GFP =  $34.2 \pm 12$ ; data are the mean  $\pm$  SEM,  $p > 0.05$ , Student's *t*-test). These results indicate that enhanced hippocampal HDAC4 expression in adulthood is sufficient to evoke depression-, but not anxiety-, like behavior.

## DISCUSSION

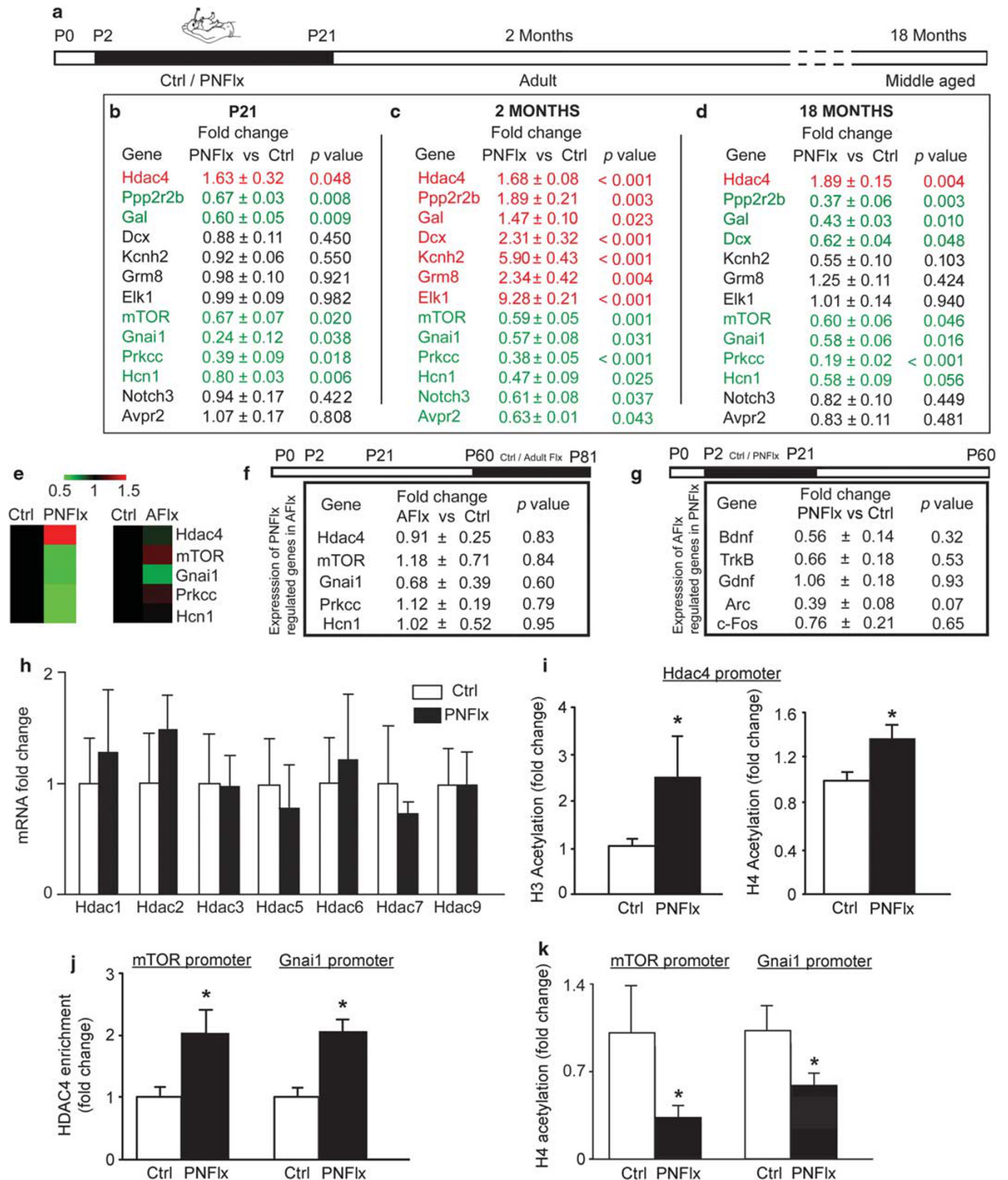
The major novel finding of our study is the identification of specific molecular signatures within the hippocampus that are associated with and contribute to the depression- and anxiety-like behaviors observed in PNFlx animals. PNFlx evoked persistent and unique gene expression changes, including a life-long increase in hippocampal *Hdac4* expression. HDAC4 enrichment at the *mTOR* and *Gnai1* promoters, accompanied by decreased activatory histone acetylation marks, support a role for HDAC4 in the downregulation of specific genes following PNFlx. Coadministration of a HDAC inhibitor prevented the effects of PNFlx on *Hdac4* and *mTOR*, and blocked the emergence of depression- and anxiety-like behavior. Further, AFlx reversed the effects of PNFlx on epigenetic modifications at the *Hdac4*, *mTOR*, and *Gnai1* promoters and normalized their hippocampal expression, accompanied by a rescue of depression- and anxiety-like behaviors. Finally, viral-mediated HDAC4 overexpression in adult hippocampus was sufficient to enhance depression- but not anxiety-like behavior. Together, our findings highlight the differing molecular signatures evoked by fluoxetine based on the temporal window of treatment, and implicate dysregulated HDAC4 in the behavioral effects of PNFlx.

Although previous studies have investigated gene expression changes following AFlx (Conti et al, 2007; Miller et al, 2008), we provide the first characterization of global transcriptional changes evoked by PNFlx. A comparison of PNFlx and AFlx transcriptomes indicates primarily nonoverlapping

**Figure 1** Postnatal fluoxetine (PNFlx) evokes persistent depression- and anxiety-like behavior and alters the hippocampal transcriptome. Shown is a schematic representation for the experimental design (a), wherein PNFlx animals were tested for juvenile play, open field test (OFT) and forced swim test (FST) in adulthood, and FST and OFT at 18 months of age. PNFlx animals showed a decline in juvenile play with reduced social grooming and pouncing (b, c) but no change in follow and chase (d). In adulthood, PNFlx animals showed depressive behavior on FST with increased immobility time (e), reduced latency to the first immobility event (f), and a trend ( $^{\#}p=0.07$ ) toward an increase in immobility events (g). PNFlx animals in adulthood showed increased anxiety-like behavior on the OFT with increased latency to approach the center (h), decline in time spent (i), and path length (j) at the center of the arena. Enhanced depression- and anxiety-like behavior on the FST (k, l) and OFT (m) persisted up to 18 months in PNFlx animals. Data are the mean  $\pm$  SEM ( $n=8-10$  animals per group for all ages). \* $p < 0.05$  as compared with controls (Student's *t*-test). Shown is a hierarchical cluster of significantly regulated (up, red; down, green) genes (n) and functional enrichment of gene ontology categories (o) in the microarray analysis performed on hippocampi (p) from 2-month-old PNFlx animals.

regulation, with enrichment of predominantly distinct gene categories. PNFlx did not influence the hippocampal expression of *Bdnf*, *TrkB*, *Gdnf*, *FGF-2*, *Arc*, and *NPY*,

which are robustly regulated by AFlx (Nibuya *et al*, 1995; Hisaoka *et al*, 2001; Miller *et al*, 2008; Christiansen *et al*, 2011) and contribute mechanistically to the neurogenic and



antidepressant effects of AFlx (Saarelainen *et al*, 2003; Sairanen *et al*, 2005; Christiansen *et al* 2011; Kohl *et al*, 2012). An important caveat is that the PNFlx and AFlx paradigms differ in treatment age and also the time after treatment when effects were assessed. With this in mind, we specifically examined whether select dysregulated genes observed in PNFlx animals in adulthood were altered right after the cessation of fluoxetine treatment. Our results revealed significant increases in *Hdac4* expression and robust decline in *mTOR*, *Prkcc*, *Hcn1*, and *Gnail* at P21, which were maintained across life in PNFlx animals. AFlx did not influence the hippocampal expression of these genes, strengthening the evidence that fluoxetine evokes distinct molecular signatures depending on the timing of treatment. A possible reason for the differential effects of fluoxetine at these two ages could be the distinct patterns of serotonin transporter expression in the postnatal vs adult brain (Narboux-Nème *et al*, 2008). Recent studies suggest that not only does the age of fluoxetine treatment administration influence the nature of behavioral and neuroplastic consequences (Iñiguez *et al*, 2010; Homberg *et al*, 2011), but that the time point at which these changes are assessed and duration following fluoxetine washout may also differentially affect outcomes.

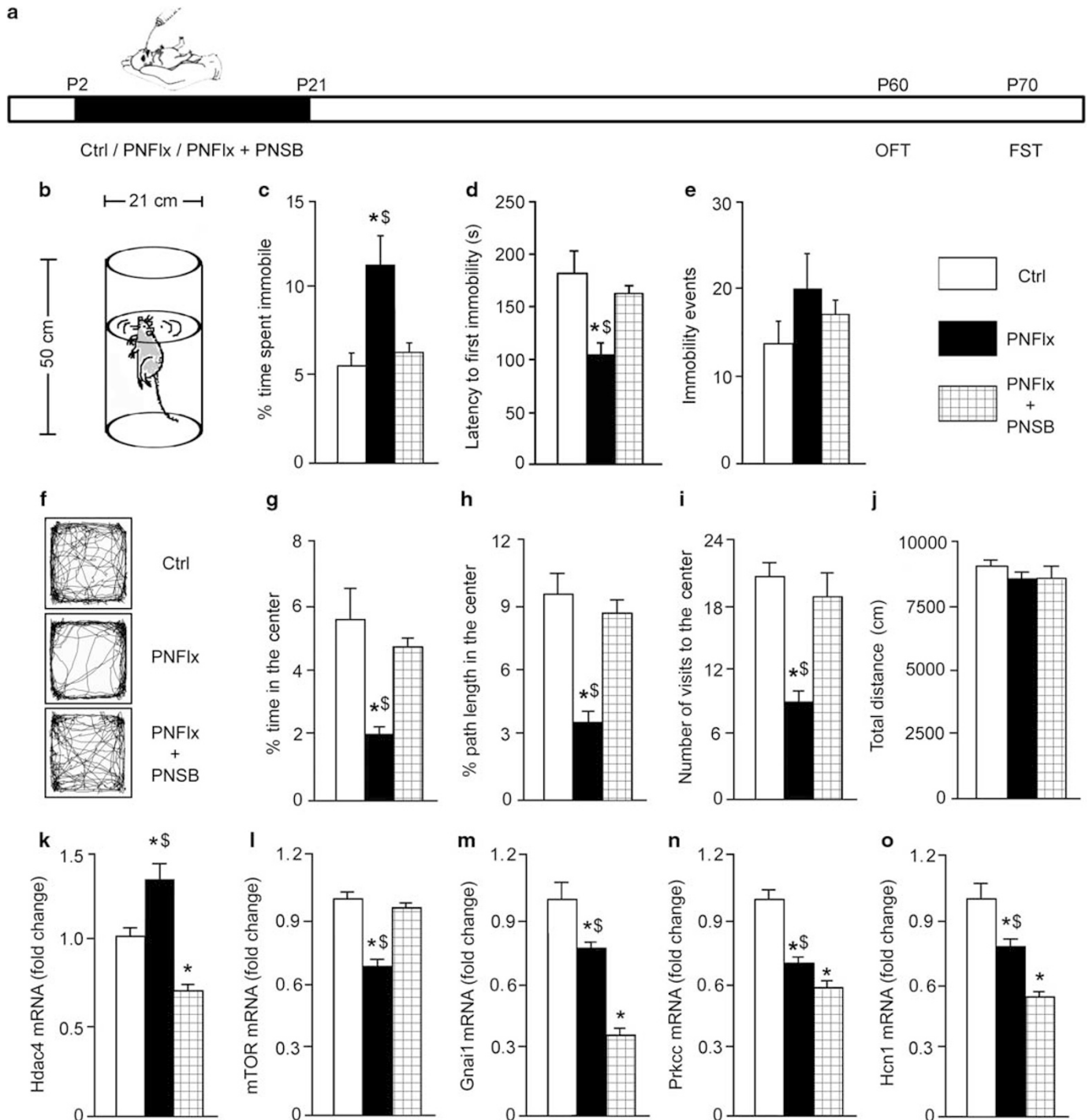
The PNFlx-evoked stable increase in *Hdac4* expression suggested that HDAC4-mediated transcriptional repression of specific genes may contribute to the emergence of behavioral abnormalities. Indeed, both concomitant postnatal HDAC inhibitor treatment and adult-onset fluoxetine retreatment, which prevented the emergence of depression- and anxiety-like behavior, were accompanied by normalization of *Hdac4* expression. Future studies are required to address the mechanisms that mediate the PNFlx-evoked sustained increase in *Hdac4* expression. *Hdac4* and *Hdac5* are highly expressed within the forebrain and influence plasticity (Kim *et al*, 2012; Sando *et al*, 2012) and mood-related behavior (Hobara *et al*, 2010; Sailaja *et al*, 2012). Recent studies show anxiolytic effects in conditional, forebrain-specific HDAC4-knockout mice (Kim *et al*, 2012). Clinical evidence indicated elevated *Hdac4* and *Hdac5* mRNA in peripheral blood cells of patients with depression (Iga *et al*, 2007; Hobara *et al*, 2010). It is interesting to speculate that HDAC4, through its histone-modifying activity, may mediate the persistent downregulation of specific genes that mechanistically contribute to the behavioral phenotypes following PNFlx.

Many more genes were downregulated following PNFlx, with DNA binding and transcriptional repression being an

enriched gene category. HDAC4 enrichment accompanied by reduced H4ac at the *mTOR* and *Gnail* promoters supports the notion that HDAC4 mediates the transcriptional repression of specific PNFlx target genes. Further, hippocampal HDAC4 overexpression also led to a decline in *mTOR* and *Gnail* expression. As a target for HDAC4-mediated repression in PNFlx animals, *mTOR* is of particular interest. PNFlx animals showed decreased hippocampal *mTOR* mRNA and protein. Normalization of PNFlx-evoked behavioral outcomes by HDAC inhibition or AFlx retreatment rescued the decreased *mTOR* expression. Dysregulated mTOR signaling is reported in neurogenetic (Gipson and Johnston, 2012), neurodevelopmental (Wang and Doering, 2013), and psychiatric disorders (Jernigan *et al*, 2011) associated with disrupted mood and social behavior. Clinical evidence indicates decreased cortical mTOR levels in depressed patients (Jernigan *et al*, 2011), whereas activation of mTOR signaling mediates the behavioral effects of rapid-action antidepressants (Li *et al*, 2010). Juvenile fluoxetine treatment (P20–34), which evokes enhanced anxiety accompanied by antidepressant-like behavior, enhanced mTOR signaling in the ventral tegmental area (Warren *et al*, 2011). The timing of fluoxetine administration may evoke distinct and region-specific effects on the mTOR pathway. Our results suggest that decreased hippocampal mTOR, likely through an HDAC4-mediated mechanism, is a stable molecular change that accompanies PNFlx-evoked depression- and anxiety-like behavior, and is normalized by treatments that rescue the behavioral effects of PNFlx.

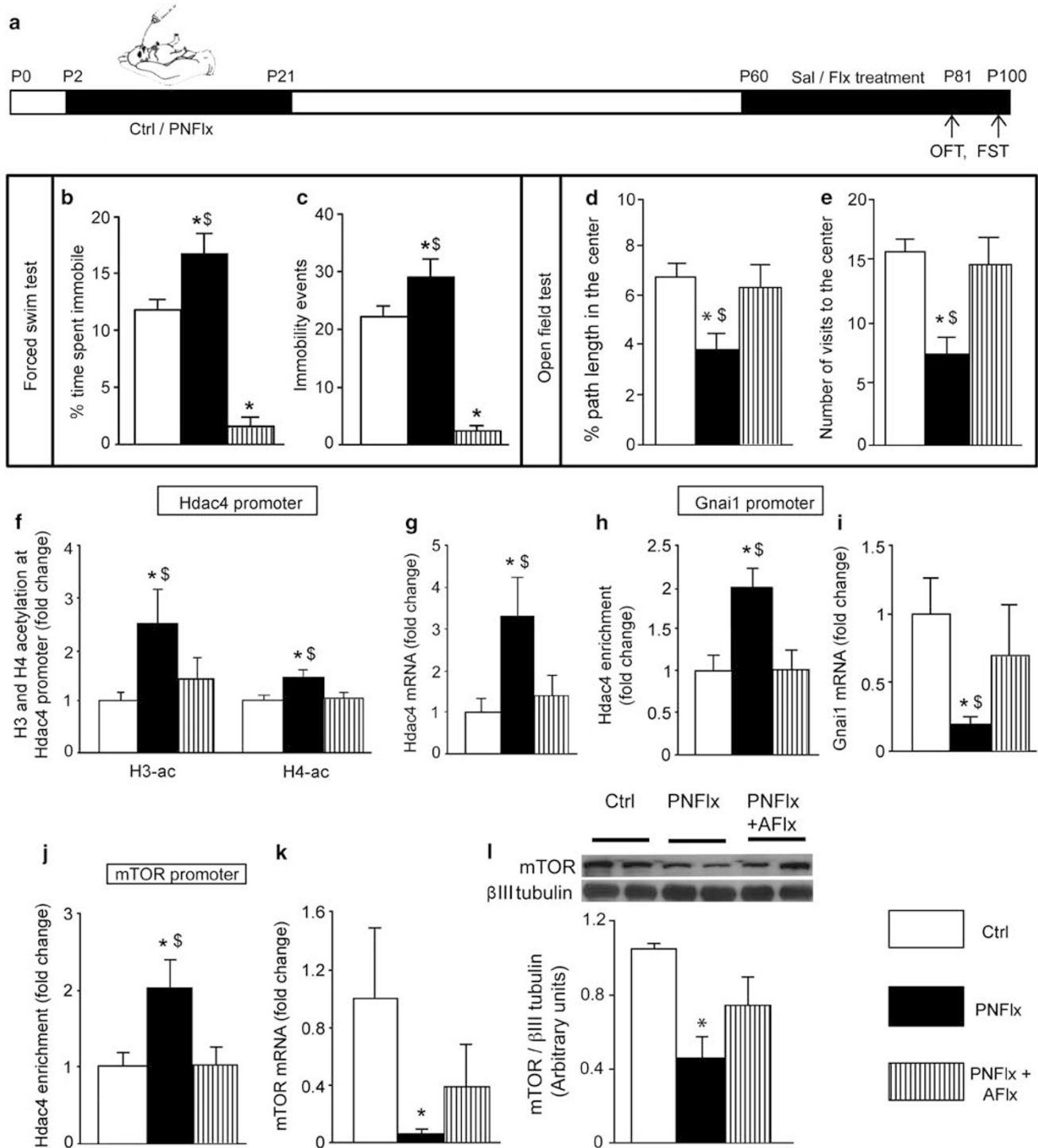
Although HDAC4 likely contributes to the transcriptional repression of *mTOR* and *Gnail*, future studies using HDAC4 ChIP-seq are required to identify the entire subset of HDAC4-regulated PNFlx target genes. Our focus on the PNFlx-mediated *Hdac4* and *mTOR* dysregulation does not preclude a role for other factors that work in concert to generate the complex PNFlx behavioral phenotype. The reversal of molecular changes (*Hdac4* and *mTOR*) by both postnatal SB and AFlx, distinct pharmacological strategies used in nonoverlapping temporal windows, suggests that rescue of specific molecular adaptations may underlie the blockade of PNFlx-evoked behaviors. A caveat to keep in mind is that SB is a broad spectrum HDAC inhibitor, which in addition to inhibiting HDAC4 activity would also perturb acetylation/deacetylation through other HDACs (Ajamian *et al*, 2004). On the basis of prior studies with valproate, which besides inhibiting HDACs evokes proteasomal

**Figure 2** Postnatal fluoxetine (PNFlx) treatment alters hippocampal gene expression and enhances histone deacetylase-4 (HDAC4) recruitment to the mammalian target of rapamycin (*mTOR*) and guanine nucleotide-binding protein alpha inhibiting 1 (*Gnail*) promoters. Shown is a schematic representation for the experimental design (a). Control (Ctrl) and PNFlx animals were killed at three different ages: P21, 2, and 18 months of age, for quantitative PCR (qPCR) analysis of hippocampal gene expression changes. Shown in the table are transcriptional changes in PNFlx animals at P21 (b), 2 months (c), and 18 months (d). qPCR analysis on an independent cohort of 2-month-old PNFlx and Ctrl animals confirmed the array-identified upregulation of *Hdac4*, *Ppp2r2b*, *Gal*, *Dcx*, *Kcnh2*, *Gm8*, *Elk1* and the downregulation of *mTOR*, *Gnail*, protein kinase C gamma (*Prkcc*), hyperpolarization-activated cyclic nucleotide-gated channel 1 (*Hcn1*), *Notch3*, *Avpr2* mRNA levels (c). The dysregulation of *Hdac4*, *mTOR*, *Gnail*, *Prkcc*, and *Hcn1* expression was observed across the lifespan in PNFlx animals (b–d). Adult fluoxetine (AFlx) treatment did not alter the hippocampal expression of *Hdac4*, *mTOR*, *Gnail*, *Hcn1*, and *Prkcc* (e, f). PNFlx treatment did not alter the expression of genes commonly upregulated by fluoxetine treatment in adulthood (AFlx) (g). PNFlx did not alter the expression of other histone deacetylases in the hippocampus (h). *Hdac4* upregulation in 2-month-old PNFlx animals was accompanied by significant increases in H3 and H4 acetylation at the *Hdac4* promoter (i). Chromatin immunoprecipitation (ChIP) analysis at 2 months of age, revealed a significant HDAC4 enrichment at the *mTOR* and *Gnail* (j) promoters in PNFlx animals. This was accompanied by significant decreases in H4 acetylation at the *mTOR* and *Gnail* promoters (k). Data are expressed as fold change of control and are the mean  $\pm$  SEM ( $n = 6$ –10 per group). \* $p < 0.05$  as compared with controls (Student's *t*-test).

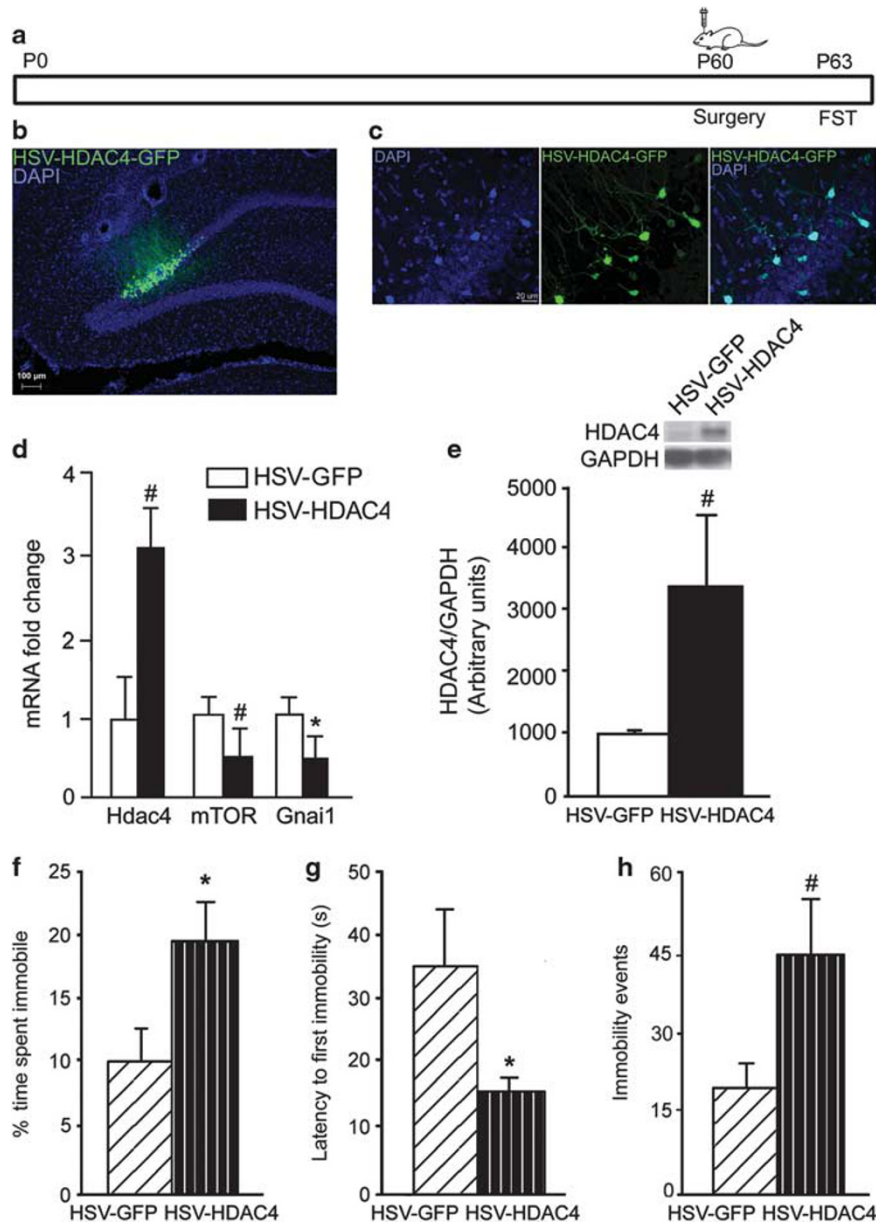


**Figure 3** Postnatal treatment with sodium butyrate (SB) prevents the emergence of postnatal fluoxetine (PNFix)-evoked molecular and behavioral changes. Shown is a schematic representation for the experimental design (a), wherein rat pups received vehicle control (Ctrl), PNFix, or fluoxetine + sodium butyrate (PNFix + PNSB) from P2 to P21 and were subjected to the forced swim test (FST) and open field test (OFT) in adulthood. PNFix animals showed a significant increase in immobility time (c) and a decreased latency to first immobility (d) on the FST (b). Coadministration of the HDAC inhibitor sodium butyrate prevented the emergence of depressive behavior on the FST (c and d). Number of immobility events remained unaltered between groups (e). Shown are representative tracks in the OFT arena from Ctrl, PNFix, and PNFix + PNSB animals (f). Coadministration of sodium butyrate prevented the anxiety-like behavior shown by PNFix animals on the OFT on measures of percent time (g), path length (h), and number of visits to the center of the arena (i). Total distance moved in the arena was unaltered across groups (j). Data are the mean  $\pm$  SEM. ( $n = 6-8$  per group,  $*p < 0.05$  as compared with Ctrl,  $^{\$}p < 0.05$  as compared with PNFix + PNSB, one-way ANOVA, the Student–Newman–Keuls *post hoc* test). Coadministration of sodium butyrate also prevented the PNFix-evoked regulation of histone deacetylase-4 (*Hdac4*; k) and mammalian target of rapamycin (*mTOR*; l), but not of guanine nucleotide-binding protein alpha inhibiting 1 (*Gnai1*; m), protein kinase C gamma (*Prkcc*; n), and hyperpolarization-activated cyclic nucleotide-gated channel 1 (*Hcn1*; o). ( $n = 6-10$  per group,  $*p < 0.05$  as compared with Ctrl,  $^{\$}p < 0.05$  as compared with PNFix + PNSB, one-way ANOVA, Student–Newman–Keul's *post hoc* test).





**Figure 4** Adult fluoxetine (AFx) treatment reverses behavioral and specific molecular signatures associated with postnatal fluoxetine (PNFlx). Shown is a schematic representation of the experimental design (a) wherein PNFlx animals in adulthood received vehicle or fluoxetine to give rise to three groups (control (Ctrl), PNFlx, and PNFlx + AFx). Retreatment in adulthood with fluoxetine rescued the depressive- and anxiety-like behavior noted on the forced swim test (FST; b, c) and open field test (OFT; d, e) in PNFlx animals. PNFlx + AFx animals did not show the PNFlx-evoked increase in immobility time (b) and number of immobility events (c), rather they exhibited antidepressant-like behavior. PNFlx + AFx animals did not show the PNFlx-evoked anxiety on the OFT on percent path length (d) and number of visits to the center of the arena (e). Further, retreatment in adulthood with fluoxetine also ameliorated the PNFlx-evoked increase in H3 and H4 acetylation at the histone deacetylase-4 (*Hdac4*) promoter (f), rescued the PNFlx-mediated increase in hippocampal *Hdac4* expression (g), normalized HDAC4 enrichment at the guanine nucleotide-binding protein alpha inhibiting 1 (*Gnai1*; h) and mammalian target of rapamycin (*mTOR*; j) promoters, and attenuated the decline in *Gnai1* (i) and *mTOR* (k) expression. Western blot analysis indicated that the PNFlx-evoked decline in mTOR levels was attenuated in the PNFlx + AFx group (l). Data are the mean  $\pm$  SEM. ( $n = 6-8$  per group for FST;  $n = 9-10$  per group for OFT;  $n = 6-10$  for ChIP/quantitative PCR (qPCR) analysis, and  $n = 4$  for western blot analysis. \* $p < 0.05$  compared with controls,  $^{\$}p < 0.05$  compared with PNFlx + AFx, one-way ANOVA, Student–Newman–Keul’s *post hoc* test).



**Figure 5** Hippocampal histone deacetylase-4 (*Hdac4*) overexpression in adulthood evokes depression-like behavior. Shown is the schematic for the experimental paradigm (a) wherein adult rats received bilateral stereotactic injections into the hippocampus of bicitronic construct HSV-HDAC4-GFP or HSV-GFP and were subjected to the forced swim test (FST). Shown are representative images for HSV-HDAC4-GFP (green) and DAPI (blue) expression in the hippocampus (b) and GFP-DAPI colocalization in the dentate gyrus (c). Quantitative PCR (qPCR) and western analysis confirmed HDAC4 overexpression with a trend noted toward significance in the qPCR ( $p = 0.07$ ; d) and western blot results ( $p = 0.062$ ; e). HDAC4 overexpression resulted in a significant decline in guanine nucleotide-binding protein alpha inhibiting 1 (*Gnai1*) expression and a strong trend ( $p = 0.051$ ) for a decline in mammalian target of rapamycin (*mTOR*) expression. HDAC4 overexpression significantly increased percent immobility time (f) and reduced latency to the first immobility event (g) in the FST. A strong trend ( $p = 0.07$ ) toward increase in the number of immobility events was noted in HDAC4-GFP-overexpressing animals (e). Data are the mean  $\pm$  SEM ( $n = 6-7$  animals per group for behavioral studies;  $n = 4$  for qPCR and western blot analysis). \* $p < 0.05$ , # $p < 0.10$  as compared with HSV-GFP (Student's *t*-test).

degradation of HDAC2 (Kramer *et al*, 2003), we speculate that a decline in overall HDAC4 levels in PNFlx + PNSB animals could contribute to the normalization of mTOR expression. Our results indicate that co-treatment with SB prevents the emergence of PNFlx-evoked depression- and anxiety-like behavior. Prior reports of SB treatment to adult animals indicate an augmentation of the behavioral effects of SSRIs (Schroeder *et al*, 2007). This suggests contrasting effects of HDAC inhibition on the

behavioral effects of fluoxetine based on the temporal window of treatment.

Our findings with AFlx retreatment of PNFlx animals indicate that interventions in adulthood can also counteract negative behavioral outcomes following PNFlx. These results corroborate prior evidence that AFlx rescues depression- and anxiety-like behavior in PNFlx animals (Karpova *et al*, 2009). AFlx-mediated rescue of behavior was accompanied by normalization of *Hdac4*, *mTOR*, and *Gnai1*

expression, and a reversal of HDAC4 enrichment at the *mTOR* and *Gnai1* promoters. AFLx itself did not influence *Hdac4*, *mTOR*, and *Gnai1* expression, raising the intriguing possibility that AFLx influences *Hdac4* expression and HDAC4-regulated targets only in the context of prior PNFlx exposure. A previous study indicates that the antidepressant imipramine downregulates HDAC5 in socially defeated animals but not in controls (Tsankova et al, 2006), further indicating that antidepressant-evoked molecular changes are context dependent. Importantly, evidence that AFLx can rescue PNFlx-induced behavioral abnormalities suggests that such changes, although potentially life long, are not immutable and can be influenced in adulthood by therapeutic strategies.

Our studies with viral-mediated hippocampal HDAC4 overexpression showed a significant increase in depression-like behavior in the FST. Given the lasting increases across the life span noted in hippocampal *Hdac4* expression following PNFlx, our results suggest that such enhanced HDAC4 may mediate effects on depression-like symptoms noted in animals with a history of PNFlx. Although we did not observe increased anxiety-like behavior following HDAC4 overexpression, our results do not allow us to rule out a role for HDAC4 in PNFlx-evoked anxiety, as early-onset changes in HDAC4 may be required for the anxiogenic effects of PNFlx. HDAC4 overexpression in adulthood in treatment-naive animals resulted in a decline in hippocampal *mTOR* and *Gnai1* levels, further lending credence to the hypothesis that HDAC4 mediates the regulation of specific PNFlx target genes.

In summary, we provide evidence of molecular signatures selectively associated with fluoxetine treatment in postnatal life, a temporal window wherein treatment evokes persistent depression- and anxiety-like behaviors. We posit that HDAC4 induction through effects on chromatin remodeling results in the establishment and maintenance of stable molecular abnormalities that contribute to the behavioral phenotypes of PNFlx. Further, we demonstrate that specific molecular changes, as well as depression- and anxiety-like behaviors, are rescued by potential therapeutic interventions in postnatal or adult life. Our results provide novel leads for the development of new approaches for treating a range of depression and anxiety syndromes, and support the building evidence that HDAC4 is a putative drug target for antidepressant therapy.

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