

Essential Role of SIRT1 Signaling in the Nucleus Accumbens in Cocaine and Morphine Action

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Sirtuins (SIRT)s, class III histone deacetylases, are well characterized for their control of cellular physiology in peripheral tissues, but their influence in brain under normal and pathological conditions remains poorly understood. Here, we establish an essential role for SIRT1 and SIRT2 in regulating behavioral responses to cocaine and morphine through actions in the nucleus accumbens (NAc), a key brain reward region. We show that chronic cocaine administration increases SIRT1 and SIRT2 expression in the mouse NAc, while chronic morphine administration induces SIRT1 expression alone, with no regulation of all other sirtuin family members observed. Drug induction of SIRT1 and SIRT2 is mediated in part at the transcriptional level via the drug-induced transcription factor Δ FosB and is associated with robust histone modifications at the *Sirt1* and *Sirt2* genes. Viral-mediated overexpression of SIRT1 or SIRT2 in the NAc enhances the rewarding effects of both cocaine and morphine. In contrast, the local knockdown of SIRT1 from the NAc of floxed *Sirt1* mice decreases drug reward. Such behavioral effects of SIRT1 occur in concert with its regulation of numerous synaptic proteins in NAc as well as with SIRT1-mediated induction of dendritic spines on NAc medium spiny neurons. These studies establish sirtuins as key mediators of the molecular and cellular plasticity induced by drugs of abuse in NAc, and of the associated behavioral adaptations, and point toward novel signaling pathways involved in drug action.

Introduction

The nucleus accumbens (NAc), central in the brain's reward circuitry, shows dramatic changes in synaptic function, neuronal morphology, and gene expression after chronic exposure to drugs of abuse. Increasing evidence suggests that aberrant epigenetic regulation is one critical factor underlying pathological responses in animal models of drug addiction and may mediate the long-lasting behavioral abnormalities that characterize addiction (Renthal and Nestler, 2008). An attractive feature of epigenetic mechanisms is the ability to rapidly respond, integrate, and process a diverse array of environmental stimuli into stable changes in chromatin structure (Kouzarides, 2007). One major form of chromatin modification is histone acetylation, catalyzed by histone acetyltransferase (HATs) and reversed by histone deacetylases (HDACs). Acetylation of specific lysine residues on histone

N-terminal tails is associated with increased gene transcription and is implicated in addiction-related phenomena (Kumar et al., 2005; Wang et al., 2010; Levin et al., 2011; Rogge and Wood, 2013). Sirtuins (SIRT)s are categorized as class III nicotinamide adenine dinucleotide (NAD)-dependent HDACs, although they deacetylate many nonhistone substrates as well. The most studied sirtuin is SIRT1, which through regulating the acetylation state of several histone and nonhistone proteins is implicated in diverse aspects of cellular physiology (Kaeberlein et al., 1999; Bellet and Sassone-Corsi, 2010; Sassone-Corsi, 2012). SIRT1 preferentially deacetylates histone H3 on lysines 9 and 14 (H3K9 and H3K14), and histone H4 on lysine 16 (H4K16; Kouzarides, 2007; Michán and Sinclair, 2007). In addition, SIRT1 has been shown to deacetylate certain nonhistone proteins, such as several transcription factors [e.g., nuclear factor- κ B (NF- κ B)], several FOXO proteins] and HATs (e.g., p300/CBP-associated factor) (Finkel et al., 2009).

Although most studies of SIRT1 to date have focused on peripheral tissues, more recent evidence suggests that the protein is also important in brain function. SIRT1 regulates neuronal differentiation and prevents neurodegeneration in several mouse models (Kim et al., 2007; Michán and Sinclair, 2007; Donmez et al., 2010). Moreover, SIRT1 is important for maintaining acquisition and consolidation of short- and long-term hippocampus-dependent memories (Gao et al., 2010; Michán et al., 2010). In a previous study (Renthal et al., 2009), using genome-wide chromatin immunoprecipitation (ChIP) followed by promoter chip

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analysis (termed “ChIP-chip”) methods, we identified SIRT1 and its closely related family member SIRT2 as targets of increased H3 acetylation in NAc after chronic cocaine administration, and verified the induction of both proteins under these conditions. We also demonstrated that systemic administration of a nonselective pharmacological activator of all sirtuins increases cocaine reward. Here, we follow up on these initial observations by establishing an essential role of SIRT1 and SIRT2 in NAc in controlling both cocaine and morphine reward by use of far more selective tools of viral-mediated gene transfer and genetic mutant mice. We also extend this work by demonstrating the actions of SIRT1 on the molecular and cellular adaptations induced in the NAc by chronic drug exposure.

Materials and Methods

Animals. C57BL/6 male mice (The Jackson Laboratory), 7–8 weeks old and weighing 25–30 g, were habituated to the animal facility 1 week before use and maintained at 23–25°C on a 12 h light/dark cycle. To induce deletion of the SIRT1 transcript in the NAc, we used mutant mice floxed for the *Sirt1* allele, which are fully backcrossed onto C57BL/6 and have been described in detail previously (Li et al., 2007). Mice were stereotaxically injected into the NAc with adeno-associated virus (AAV) vectors (serotype 2) expressing GFP or Cre-GFP between the age of 7 and 10 weeks. All animals had access to food and water *ad libitum*. All experiments were approved by the Animal Care and Use Committee at Mount Sinai Medical Center.

Drugs. Cocaine-HCl and morphine-SO₄ were obtained from Sigma-Aldrich or the National Institute on Drug Abuse (Bethesda, MD). Drugs were given as daily intraperitoneal injections for 7 d at 20 mg/kg. Acute studies consisted of 6 d of saline injections followed by a single drug injection on the seventh day. Control groups received 7 d of saline injections. Animals were analyzed at varying times after the last injection.

RNA isolation and PCR. Bilateral 14 gauge NAc punch dissections were obtained from 1-mm-thick coronal brain sections and were frozen immediately on dry ice. Dissections samples were thawed and processed in TriZol (Invitrogen) according to manufacturer guidelines. RNA was isolated and purified using RNeasy Micro columns (Qiagen). We confirmed purity of our samples by spectroscopy at 260/280 and 260/230 > 1.8. RNA was reverse transcribed to cDNA using iScript Kit (Bio-Rad). Quantitative PCR (qPCR) was performed using an Applied Biosystems 7500 system. Reactions were run in triplicate and analyzed using the $\Delta\Delta C_t$ method (Livak and Schmittgen, 2001).

Chromatin immunoprecipitation. ChIP was performed for Δ FosB as described previously (Kumar et al., 2005), with minor modifications. Briefly, for each ChIP, anterior and posterior bilateral 14 gauge NAc punches were pooled (five animals, 20 NAc punches per sample). Punches were fixed for 10 min with 1% formaldehyde and then quenched with 2 M glycine for 5 min. Samples were homogenized using a desktop sonicator at low settings (amplitude 40%) two times for 7 s on ice. Next, samples were sheared using a Diagenode Bioruptor XL at 4°C at high sonication intensity for 30 s on/30 s off for 25 min, followed by 25 min of rest and an additional 25 min of sonication. Fragment size of 250–1000 bp was verified with an Agilent Bioanalyzer. Before sonication of samples, magnetic sheep anti-rabbit or anti-mouse beads (Invitrogen) were prepared with the respective antibody of interest at 4°C overnight on a rotator. Following washing of the magnetic bead/antibody complex, 7.5 mg (magnetic bead/antibody complex) was added to 400 μ l of sheared chromatin for histone ChIPs, or 1000 μ l for Δ FosB or other nonhistone protein ChIPs, for 16 h at 4°C. Eighty microliters of each sample of sheared chromatin was used as an input control. Samples were washed with LiCl and Tris-EDTA buffers. Reverse cross-linking was performed at 65°C overnight, and proteins were removed with proteinase K (Invitrogen). DNA was purified using a DNA purification kit (Qiagen). Additionally, a normal IgG control was performed to test for nonspecific binding.

ChIP-Seq. Animals were treated with chronic cocaine administration (20 mg/kg) daily for 7 d, and 24 h later NAc tissue pooled from approx-

imately five mice was processed for ChIP as described previously (Renthal et al., 2009; Maze et al., 2010). Antibodies were all ChIP grade from Abcam. Approximately 10 ng of input DNA or pull-down DNA was used for the preparation of sequencing libraries following the instructions of the Illumina ChIP-Seq Sample Prep Kit (catalog #IP-102-1001). In brief, DNA fragment overhangs were converted into phosphorylated blunt ends using T4 DNA polymerase, Klenow polymerase, and T4 polynucleotide kinase. An “A” base was then added to the DNA fragments to enable ligation to the adapters, which have a single “T” overhang. The libraries were analyzed on a 2% agarose gel, and size was selected between 175 and 300 bp. Gel-extracted DNA was further enriched by PCR and run on a bioanalyzer to validate size distribution and concentration. All libraries were sequenced on Illumina Hi-Seq 2000 machines at Mount Sinai Hospital and were analyzed according to published methods (Maze et al., 2011; Shen et al., 2013).

Immunoblotting. Mice were given a lethal dose of chloral hydrate, then sequentially perfused with PBS and 4% paraformaldehyde. Brains were incubated in a 30% sucrose solution overnight at 4°C and sectioned at 35 μ m on a microtome (Leica). Free-floating sections were washed with PBS, then blocked (donkey serum), and later incubated with anti-GFP (rabbit polyclonal, 1:200; Invitrogen) and/or anti-SIRT1 (rabbit polyclonal 1:500; Millipore). After 4°C overnight incubation with constant agitation, sections were washed and incubated with Cy2 and/or Cy3 fluorescent antibodies for 2 h at room temperature, after which they were washed, step grade ethanol dehydrated, and mounted with DPX. Sections were then analyzed using confocal microscopy with a Carl Zeiss 560LSM instrument at various magnifications. Immunoblotting was performed using standard procedures on NAc punch dissections from individual animals (Maze et al., 2010). Final blots were developed by chemiluminescence analysis using supersignal dura ECL (Pierce Biotechnology/Thermo Fisher Scientific). Images were quantified using densitometry (ImageJ, NIH), and samples were normalized using GAPDH or β -tubulin.

Viral-mediated gene transfer. Mice were anesthetized with a ketamine/xylazine cocktail (ketamine 100 mg/kg and xylazine 10 mg/kg) and prepared for stereotaxic surgery. Thirty-three gauge syringe needles (Hamilton) were used to bilaterally infuse 0.5 μ l of virus into NAc at a rate of 0.1 μ l/min at 1.6 mm anteroposterior, +1.5 mm lateral, and 4.4 mm dorsoventral from bregma. We used bicistronic p1005 herpes simplex virus (HSV) vectors expressing GFP alone or GFP plus SIRT1, SIRT2, or a CreGFP fusion protein. In this system, GFP expression is driven by a cytomegalovirus promoter, whereas the select gene of interest is driven by the IE4/5 promoter (Maze et al., 2010). In some experiments, we used AAV vectors serotype 2 expressing GFP or CreGFP as described previously (Maze et al., 2010).

Conditioned place preference. Conditioned place preference (CPP) was performed as described previously (Maze et al., 2010). Briefly, animals were first prescreened in a photobeam-monitored box to detect any baseline bias to the two chambers of the CPP apparatus. Mice that showed a statistically significant preference to one of the two chambers were removed from further analysis (<10% of all animals). Mice were then divided into control and experimental groups with equivalent pretest scores. During the training phase, animals were injected with saline in the morning and placed in one chamber for 30 min, then were injected with cocaine (5–15 mg/kg, i.p.) and placed in the other chamber in the afternoon. This training regimen was conducted for 2 d followed by a test day in which mice were placed back into the apparatus without treatment for 20 min and evaluated for chamber preference. For all groups, baseline locomotion in response to saline was assessed. CPP scores were calculated by subtracting time spent in the saline-paired chamber from time spent in the cocaine-paired chamber. Please note that different drug doses were used depending on the experimental conditions: lower doses were used for experiments in which the manipulation was predicted to enhance CPP to avoid ceiling effects, and higher doses were used for experiments in which the manipulation was predicted to reduce CPP to avoid floor effects.

Locomotor sensitization. Locomotor sensitization was measured per published protocols (Lobo et al., 2010) with minor modifications. Activity was assessed in the *x*- and *y*-planes for horizontal ambulations in a 75

cm² chamber using EthoVision XT. On days 1–7, mice were first habituated to the apparatus for 5 min then injected intraperitoneally with saline, and 10 min later they were injected with saline or cocaine (7.5 mg/kg) and analyzed for 30 min. Seven days after the last saline/cocaine injection, they were given a challenge injection of cocaine and again tested for locomotor activity.

Statistical analysis. One- and two-way ANOVAs were performed to determine significance for conditions in which there were more than two groups or two factors. Unpaired Student's *t* test with a two-tailed *p* value were used for other comparisons including qPCR, CPP, and Western blotting, comparing HSV-GFP to HSV-Cre in *Sirt1* floxed mice, and ChIP experiments. All values included in the figure legends represent means ± SEM.

Results

Drugs of abuse induce SIRT1 and SIRT2 expression in the NAc

As a first step to explore the role of sirtuins in drug action, we analyzed mRNA expression levels of all SIRT family members in mouse NAc after acute or chronic (7 d) exposure to cocaine or morphine, with animals analyzed 24 h after the last drug dose. As shown in Figure 1A, chronic cocaine induces *Sirt1* and *Sirt2* mRNA levels in NAc, whereas chronic morphine induces *Sirt1* only, with no changes seen in all other sirtuins. These effects were regionally specific as we did not observe any significant changes in the dorsal striatum (data not shown). To define the temporal pattern of SIRT regulation, we collected NAc from animals treated chronically with cocaine or morphine at several time points after the last drug dose. While the chronic cocaine-induced increase in *Sirt1* and *Sirt2* mRNA levels were apparent at 4 and 24 h with no change seen at the 1 h time point (Fig. 1B), the chronic morphine-induced increase in *Sirt1* was apparent at 24 h only (Fig. 1C). We also found that the cocaine- and morphine-induced increase in *Sirt1* expression, but not *Sirt2* expression, was relatively long lived: increased *Sirt1* mRNA levels persisted through 5 d of withdrawal and reverted to control levels by day 10 (Fig. 1B,C). Western blot analysis demonstrated that similar drug regulation was seen at the level of SIRT1 and SIRT2 protein. At the 24 h time point, chronic administration of cocaine or morphine increased levels of SIRT1 protein in NAc, whereas only cocaine increased levels of SIRT2 protein (Fig. 1D,E). In contrast, we observed no change in *Sirt1* or *Sirt2* mRNA or protein levels at either 4 or 24 h after single doses of cocaine or morphine (Fig. 1F,G), indicating that repeated exposure is required for induction of these genes.

Next, to gain insight into whether drug regulation of SIRT expression in NAc is achieved at the transcriptional level, we took advantage of large ChIP-Seq datasets for several chromatin marks in NAc of mice treated chronically with cocaine or saline and examined 24 h after the last treatment. Mining these existing data for changes that occur with cocaine at the *Sirt1* gene, we found changes consistent with transcriptional activation: increased binding of a mark associated with active gene expression such as trimethylation of lysine 4 of H3 (H3K4me3), and no changes in marks associated with gene repression such as dimethylation and trimethylation of lysine 9 of H3 (H3K9me2 and H3K9me3; Fig. 1H). These observations support the hypothesis that cocaine induction of SIRT1 is mediated via transcriptional mechanisms.

Mechanisms underlying SIRT1 and SIRT2 induction in the NAc

Based on these findings, we studied the upstream transcriptional mechanisms mediating cocaine and morphine induction of SIRT1 and SIRT2. Δ FosB, a Fos family transcription factor that

binds to activator protein 1 (AP1) sites in responsive genes, is well characterized for its role in NAc in addiction-related behaviors (Nestler, 2008; Robison and Nestler, 2011). Δ FosB is a highly stable protein and, once induced, persists in NAc for several weeks. A previous ChIP-chip study showed significant enrichment of Δ FosB binding to the *Sirt2* gene promoter in NAc of mice treated chronically with cocaine, with no change apparent at the *Sirt1* gene (Renthal et al., 2009). However, ChIP-chip is known to lack quantitative power. Thus, to follow up these initial observations, we performed quantitative ChIP (qChIP) and found robust (twofold to threefold) increased binding of Δ FosB to both the *Sirt1* and *Sirt2* promoters in NAc after chronic treatment with cocaine, with similar increases in Δ FosB binding seen for both genes after chronic morphine administration (Fig. 2A). The observation that chronic morphine increases Δ FosB binding to the *Sirt2* promoter, even though it does not increase *Sirt2* expression, indicates that other mechanisms are involved (see Discussion).

Δ FosB acts as a transcriptional activator or repressor depending on the gene to which it binds (McClung and Nestler, 2003; Kumar et al., 2005; Renthal et al., 2008, 2009). To determine the transcriptional effect of Δ FosB on the *Sirt1* gene, we performed a series of studies using AAV or HSV vectors to overexpress either Δ FosB or a truncated JunD protein (termed " Δ JunD"), which antagonizes AP1-mediated transcription (Peakman et al., 2003; Winstanley et al., 2007; Nestler, 2008). First, using AAV vectors to bilaterally overexpress Δ FosB in mouse NAc for 3 weeks (Fig. 2B), we observed a selective and significant increase in mRNA levels of *Sirt1* and *Sirt2* (Fig. 2C). Such induction of *Sirt1* by AAV- Δ FosB was associated with increased binding of Δ FosB to its gene promoter (Fig. 2D), indicating that virally expressed Δ FosB is functionally active. In contrast, we saw no effect of AAV- Δ FosB on other sirtuin genes (Fig. 2E). Additionally, Δ FosB induction of *Sirt1* requires sustained expression of Δ FosB, since Δ FosB overexpression with HSV vectors, which maximally express transgenes for days 1–4 only (Barrot et al., 2002), did not alter *Sirt1* levels (Fig. 2C). Next, we overexpressed Δ JunD in NAc using AAV vectors and found that this manipulation blocked the ability of cocaine to induce *Sirt1* in this brain region (Fig. 2F). Interestingly, Δ JunD did not block the binding of endogenous Δ FosB to the *Sirt1* promoter (Fig. 2D), consistent with its known ability to dimerize with Δ FosB and bind to AP1 elements in responsive genes, but disrupt transcriptional activation due to the lack of its N-terminal transactivation domain (Peakman et al., 2003; Winstanley et al., 2007; Nestler, 2008). Together, these studies demonstrate that Δ FosB functions as a transcriptional activator at the *Sirt1* and *Sirt2* genes.

In addition to transcriptional mechanisms, we identified indirect evidence for post-transcriptional and post-translational regulation of SIRT1 in NAc by chronic cocaine and morphine as well. In cell culture systems, SIRT1 expression is enhanced by the RNA binding protein Hu antigen R (HuR; also known as ELAV1), a member of the embryonic lethal abnormal vision family of mRNA binding proteins. HuR binds to the 3' UTR of *Sirt1* mRNA in cultured cells and increases its half-life by >8 h (Sasaki et al., 2006; Abdelmohsen et al., 2007). Interestingly, we found that chronic cocaine and chronic morphine administration increase levels of HuR mRNA and protein in the mouse NAc (Fig. 3A,B). There is also evidence from cell culture systems that SIRT1 is modified by sumoylation—a post-translational modification involving the addition of small ubiquitin-related modifiers (SUMOs) to lysine residues that is implicated in the regulation of several transcriptional repressors (Bossis and Melchior, 2006). SIRT1 is sumoylated at lysine 734 by

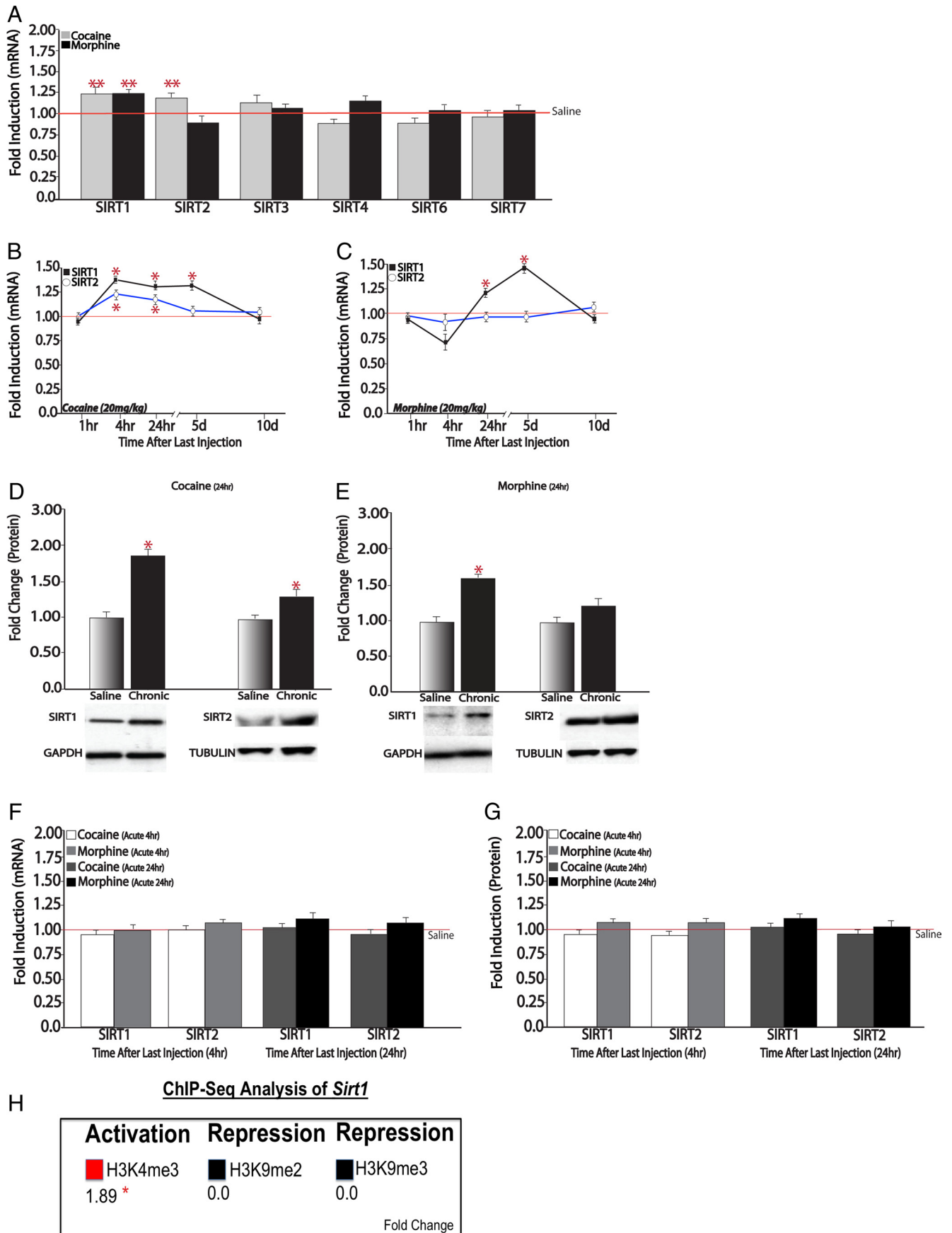


Figure 1. Drugs of abuse induce SIRT expression in NAc. **A**, qPCR analysis of NAc from chronic (7 d) cocaine-treated mice (20 mg/kg, i.p.) and chronic (7 d) morphine-treated mice (20 mg/kg, i.p.), analyzed 24 h after the last drug dose, revealed increased levels of *Sirt1* mRNA after cocaine ($n = 9-10$) or morphine ($n = 8$) administration. *Sirt2* mRNA was increased by (Figure legend continues.)

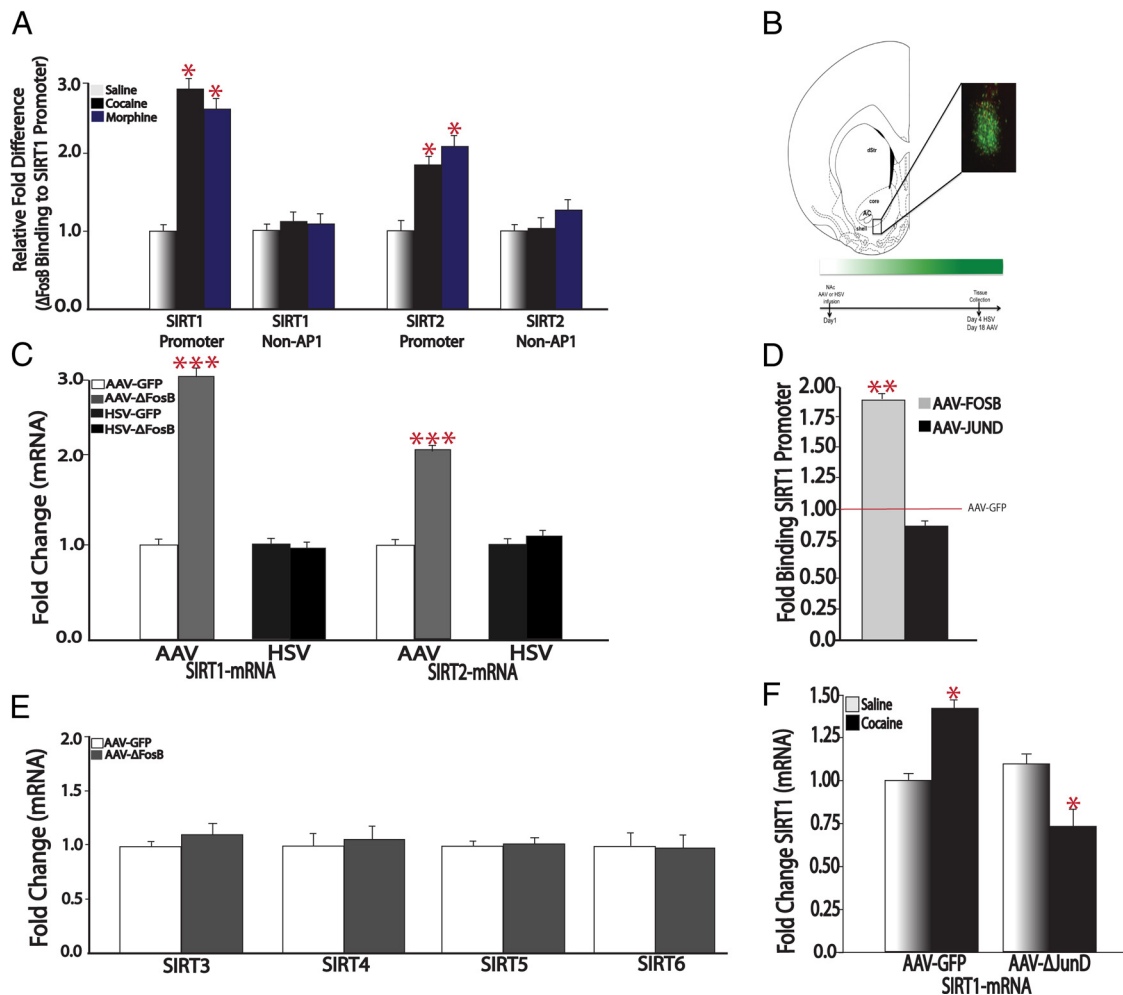


Figure 2. Role of Δ FosB in *Sirt1* and *Sirt2* induction in NAc by cocaine and morphine. **A**, qChIP reveals increased binding of Δ FosB to the *Sirt1* and *Sirt2* gene promoters in the NAc of mice treated for 7 d with cocaine or morphine (20 mg/kg, i.p.) analyzed 24 h after the last injection ($*p < 0.05$; $n = 6-8$). **B**, Representative image of GFP coexpressed with transgene in NAc. The area of infection was limited to the dorsomedial region of NAc including both shell and core. Time course of viral vector expression: AAV reaches maximal expression 10–14 d postinfusion and is maintained thereafter, consistent with previous studies (Zachariou et al., 2006; Winstanley et al., 2007). **C**, Sustained AAV-mediated overexpression of Δ FosB in NAc induces *Sirt1* mRNA ($n = 7-8$) and *Sirt2* mRNA ($n = 7$), whereas shorter-term HSV-mediated overexpression does not ($n = 7-8$). **D**, AAV- Δ FosB increases Δ FosB binding to the *Sirt1* promoter ($n = 7$), while AAV- Δ JunD had no effect ($n = 7$). **E**, AAV- Δ FosB had no effect on other sirtuins ($n = 7-8$). **F**, Overexpression of Δ JunD blocks cocaine's induction of *Sirt1* ($n = 5-9$). $*p < 0.05$, $**p < 0.01$, $***p < 0.001$.

SUMO-1, which results in an increase in SIRT1's deacetylase activity (Yang et al., 2007). We found that chronic cocaine and chronic morphine administration induce both mRNA and protein levels of SUMO-1 in the mouse NAc (Fig. 3C,D). These findings raise the possibility that drugs of abuse might also regulate SIRT1 mRNA translation and catalytic activity to

promote its function, possibilities that now require direct experimentation.

SIRT1 and SIRT2 in the NAc increase behavioral responses to cocaine and morphine

We previously implicated sirtuins in behavioral responses to cocaine by use of broadly acting and nonspecific pharmacological agents, some of which were given systemically and hence affected sirtuin activity in all tissues and brain regions (Renfahl et al., 2009). To directly test the specific involvement of SIRT1 and SIRT2 in the NAc in behavioral responses to cocaine and morphine, we used HSV vectors to overexpress SIRT1 or SIRT2 in mouse NAc and assessed CPP, an indirect measure of reward, to these drugs. First, immunohistochemical analysis confirmed the targeting of medium spiny neurons in NAc after bilateral infusion of HSV vectors; expression was circumscribed within the dorsomedial part of NAc including portions of both shell and core (Fig. 4A). To further validate the HSV-SIRT1 vector, we assessed acetylation levels of several histone and nonhistone substrates for SIRT1 in other tissues (Finkel et al., 2009; Fig. 4B).

←

(Figure legend continued.) cocaine only ($n = 7-9$). *Sirt3* to *Sirt7* were not regulated by either drug of abuse ($n = 7-10$). **B**, Chronic cocaine increases *Sirt1* levels 4 h ($n = 6$), 24 h ($n = 9-10$), and 5 d ($n = 8$) after the last 7 d injection, with no change seen at 1 h or 10 d ($n = 6-8$). A similar pattern holds for *Sirt2*, with small but significant induction apparent at 4 and 24 h ($n = 8-9$). **C**, Chronic morphine increases *Sirt1* levels at 24 h and 5 d ($n = 8$), with no regulation seen for *Sirt2* ($n = 8$). **D**, **E**, Western blotting reveals that chronic cocaine or chronic morphine administration increased SIRT1 and chronic cocaine administration increased SIRT2 protein levels at the 24 h time point ($n = 5-10$). **F**, **G**, A single dose of cocaine or morphine had no effect on *Sirt1* or *Sirt2* mRNA levels (**F**) or protein levels (**G**) at either 4 or 24 h ($n = 8-10$). **H**, ChIP-Seq analysis of chromatin marks in NAc after chronic cocaine exposure reveals increased binding of marks associated with gene activation (e.g., H3K4me3), with no changes in binding of repressive marks (H3K9me2 and H3K9me3). $*p < 0.05$, $**p < 0.01$.

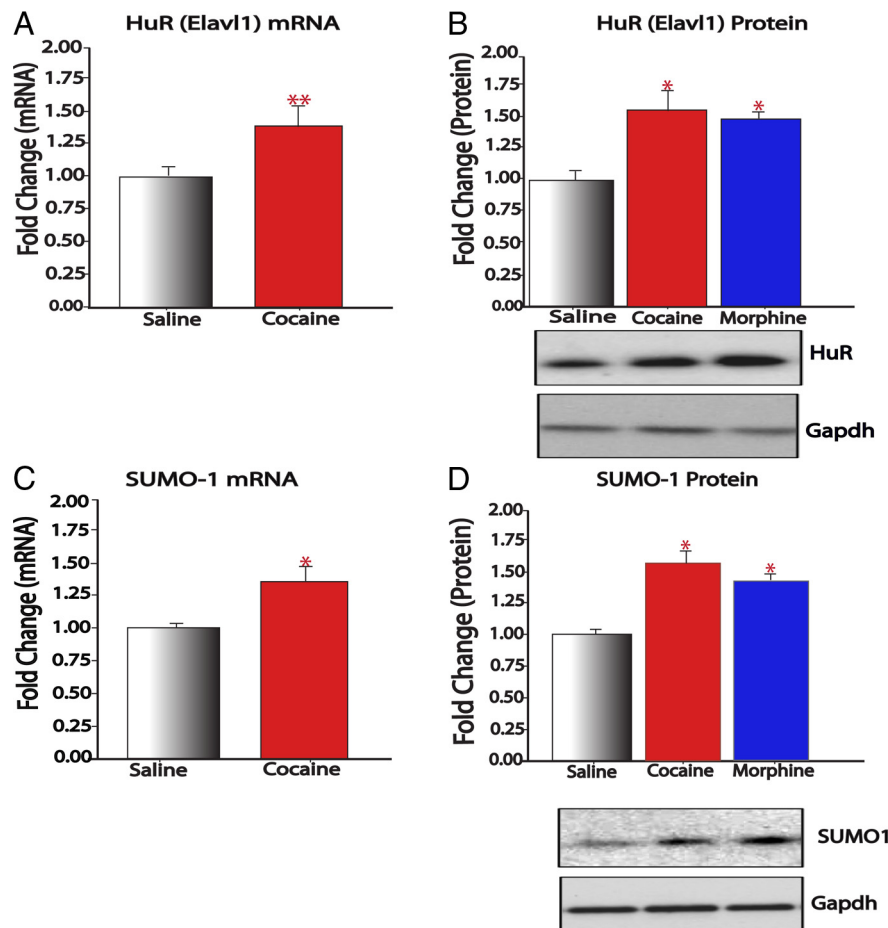


Figure 3. Regulation of post-transcriptional and post-translational modulators of SIRT1. **A, B**, Chronic (7 d) cocaine increases both HuR transcript levels (**A**; $n = 6-8$) and protein levels (**B**; $n = 7$) in NAc. A similar increase in HuR protein is induced in NAc by chronic morphine ($n = 8-10$; **B**). **C, D**, Cocaine also increases SUMO-1 transcript levels (**C**; $n = 6-8$) and protein levels (**D**; $n = 7-8$). Chronic morphine likewise induces SUMO-1 protein in NAc ($n = 8-10$; **D**). * $p < 0.05$, ** $p < 0.01$.

HSV-SIRT1, which increased SIRT1 protein levels in NAc by approximately twofold, decreased levels of acetyl-H4K16, acetyl-p65 (an NF- κ B subunit), and acetyl-FKRH (a FOXO transcription factor), with a trend for decreased global levels of pan-acetyl-H3. In contrast, levels of other SIRT substrates seen in other systems, including acetyl-H3K9, acetyl-H3K56, and acetyl-p53, were unaffected in NAc by SIRT1 overexpression or in some cases showed changes in the opposite direction. These observations support the notion that SIRT might target a partly different set of proteins in NAc compared with other brain regions or peripheral tissues. To determine whether HSV-mediated overexpression of SIRT1 is sufficient to regulate genes known to play a role in synaptic plasticity in the NAc, we treated animals intra-NAc with HSV-GFP or HSV-SIRT1 and assessed gene expression by qPCR 3 d later, which corresponds to the first day of behavioral testing in the cocaine and morphine CPP paradigm. We observed significant increases in mRNA expression of *Bdnf* (brain-derived neurotrophic factor), *Creb1* (cAMP response element binding protein-1), *Cbp* (CREB binding protein), *Cdk5* (cyclin-dependent kinase-5), *Syn2* (synapsin 2), and *Gria1* (the GluA1 AMPA glutamate receptor subunit) in the NAc of mice overexpressing HSV-SIRT1 relative to the HSV-GFP control group (Fig. 4C).

Having validated the vectors, we proceeded with CPP studies, using an unbiased paradigm in which animals do not prefer a chamber at baseline (Fig. 5A). HSV-SIRT1 increased CPP for

cocaine relative to the HSV-GFP control group in a dose-dependent fashion (Fig. 5B). Likewise, animals injected with HSV-SIRT1 showed a dose-dependent increase in morphine CPP (Fig. 5C). We observed similar increases in cocaine and morphine CPP when SIRT2 was overexpressed (Fig. 5D, E). We also explored the effect of SIRT1 on cocaine-induced locomotor sensitization. The acute locomotor-activating effects of cocaine increase with repeated drug exposure; such sensitization is thought to reflect changes in synaptic plasticity and gene expression that contribute to aspects of drug addiction (Robinson and Berridge, 2008). Mice injected with HSV-SIRT1 displayed an enhanced locomotor response to the initial injection of cocaine relative to the HSV-GFP control group (Fig. 5F, G). This effect was specific to cocaine, and was not the result of injection-related stress, because HSV-SIRT1 animals injected with saline did not exhibit an increase in locomotor activity (data not shown). This effect on cocaine responses was transient, as no further enhancement in locomotor sensitization from days 2 to 7 was observed relative to the control group (Fig. 5F). However, after 7 d of withdrawal, HSV-SIRT1 mice displayed a dramatic enhancement in locomotor sensitization in response to a challenge dose of cocaine when compared with HSV-GFP controls (Fig. 5F, G). These findings further substantiate that SIRT1 induction in NAc promotes behavioral responses to cocaine,

and show that even a transient increase in SIRT1 expression, mediated by HSV vectors, can lead to enhanced behavioral responses that persist long after transgene expression reverts to control levels.

To gain insight into the mechanism underlying this sustained effect of HSV-SIRT1, we performed qChIP for SIRT1 at several representative genes in the NAc of mice that had received injections of HSV-GFP or HSV-SIRT1 into this brain region 14 d earlier. We found a significant increase in SIRT1 binding at the *Bdnf* and *Cbp* genes, and trends for increases at the *Cdk5* and *Syn2* genes (Fig. 4D). These data indicate that a transient increase in SIRT1 levels can lead to more sustained increases in SIRT1 binding to certain target genes, providing one possible mechanism for the sustained behavioral effects observed with HSV-SIRT1.

To investigate whether SIRT1 in NAc is required for normal behavioral responses to drugs of abuse, we injected an AAV vector expressing Cre recombinase or GFP alone specifically into NAc of mice homozygous for a loxP-flanked *Sirt1* gene (Li et al., 2007). Eighteen days after AAV injection (when transgene expression is maximal), SIRT1 knockdown was confirmed by immunohistochemistry (Fig. 5H). Using this approach, we observed that animals injected with AAV-Cre displayed decreased preference for cocaine and for morphine at drug doses that produce robust CPP in AAV-GFP control mice (Fig. 5I, J). These

data demonstrate that SIRT1 activity in NAc is necessary for enhanced cocaine and morphine reward.

SIRT1 regulates dendritic spine density in the NAc

The findings that SIRT1 is crucial for reward-related behavior and can regulate plasticity-related genes in the NAc prompted us to investigate the possibility that SIRT1 may also regulate dendritic plasticity in this brain region. A large literature has shown that chronic cocaine administration increases dendritic spine density of NAc medium spiny neurons and that this structural plasticity is associated with enhanced behavioral sensitivity to cocaine (Robinson and Kolb, 2004; Kalivas, 2009; Russo et al., 2010). Here, we first showed that chronic treatment with cocaine increased the expression of several genes that have been implicated previously in regulating dendritic morphology and synaptic plasticity in NAc, such as *Bdnf*, *Creb1*, *Cbp*, and *Cdk5* (Fig. 6A). As previously reported using similar treatment conditions (Russo et al., 2009; LaPlant et al., 2010; Maze et al., 2010), we observed a corresponding increase in dendritic spine density of NAc medium spiny neurons of mice treated with chronic cocaine administration (Fig. 6B). HSV-mediated overexpression of SIRT1 in this brain region was sufficient to mimic the effects of cocaine by increasing NAc dendritic spine density (Fig. 6C). These results suggest that SIRT1 signaling may be one mechanism underlying cocaine-induced increases in NAc dendritic morphology and expression of plasticity-related genes.

Discussion

Drugs of abuse regulate SIRT1 and SIRT2 expression in the NAc

Results of the present study demonstrate an essential role of SIRT1 and SIRT2, class III HDACs, in the actions of drugs of abuse on the NAc reward circuit. Our findings show that drugs of abuse specifically regulate SIRT1 and SIRT2 expression in NAc in a temporally defined manner. While chronic cocaine administration induced both proteins in this brain region, chronic morphine administration induced SIRT1 only. These effects were specific to repeated administration of the drugs, as we did not observe changes with acute cocaine or morphine administration. Importantly, these mRNA changes occurred concomitantly with equivalent increases in SIRT1 and SIRT2 protein. Such induction of the SIRTs was associated with chromatin changes at their gene promoters normally indicative of gene activation. Thus, ChIP-Seq analysis for several types of prominent histone modifications indicated that chronic cocaine administration in-

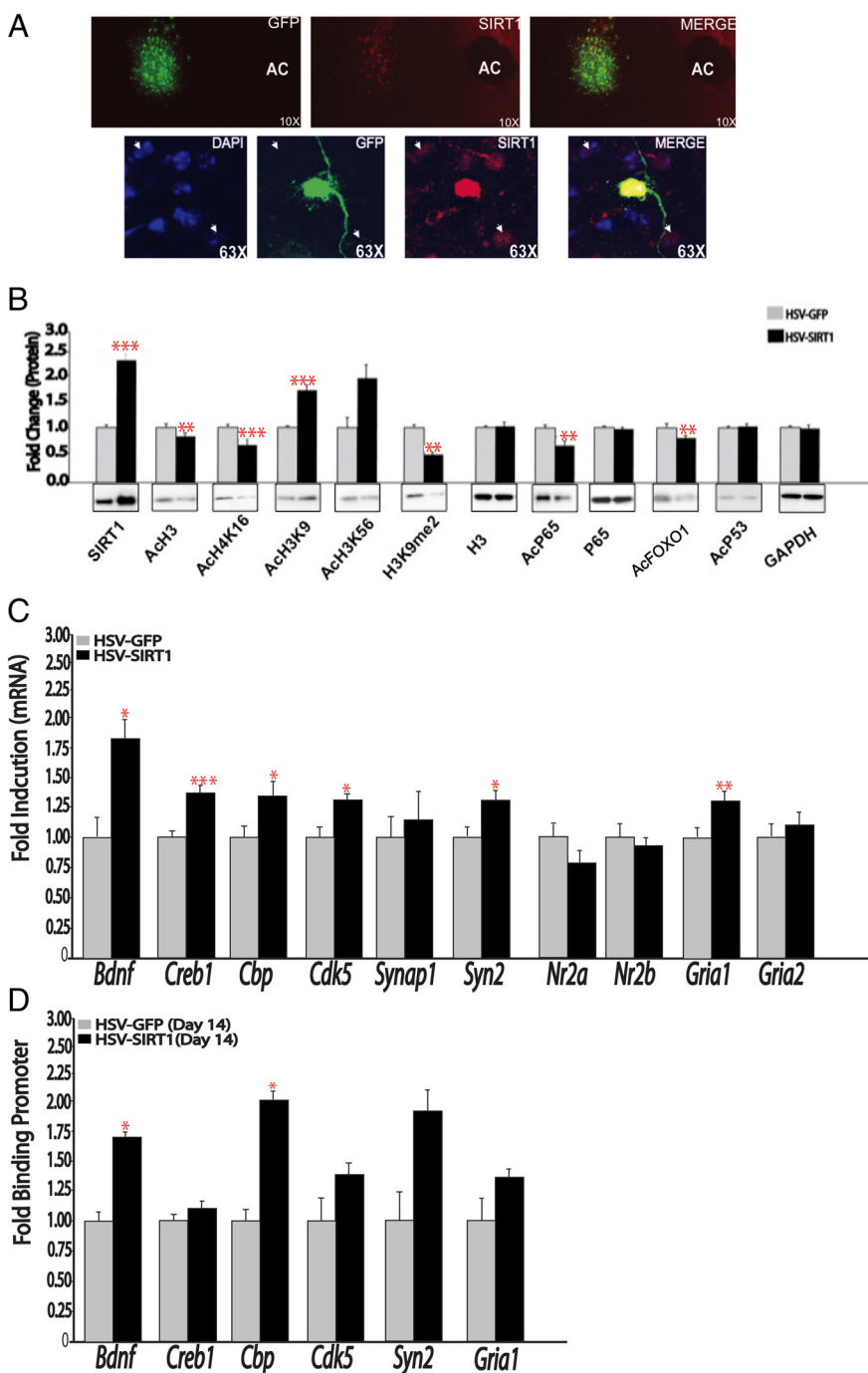


Figure 4. HSV-SIRT1 regulates synaptic plasticity genes. **A**, Double-labeling immunohistochemistry shows robust induction of SIRT1 (red) in neurons infected with HSV-SIRT1 4 d after bilateral HSV vector injections into NAc. **B**, Western blotting reveals that HSV-mediated SIRT1 overexpression in NAc deacetylates certain known histone and nonhistone substrates of SIRT1, but not others ($n = 5-11$). **C**, HSV-SIRT1 regulates the expression of certain plasticity-related genes in the NAc. qPCR analysis of mouse NAc overexpressing GFP or SIRT1 ($n = 6-8$). **D**, qChIP for *Sirt1* in the NAc of mice injected with HSV-GFP or HSV-SIRT1 and assessed for *Sirt1* binding at the promoter of several plasticity-related genes 14 d later ($n = 3-5$). * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

duces an increase in H3K4me3 binding at the *Sirt1* promoter, with no changes seen in repressive histone methylation (H3K9me2 and H3K9me3). These data extend earlier ChIP-chip findings, which showed increased acetylated H3 binding on both the *Sirt1* and *Sirt2* gene promoters as well (Renthal et al., 2009). These data indicate that chronic exposure to drugs of abuse induces SIRT1 (and, for cocaine, SIRT2 as well) in NAc via the transcriptional activation of their genes. Our findings thus add these sirtuins to several other

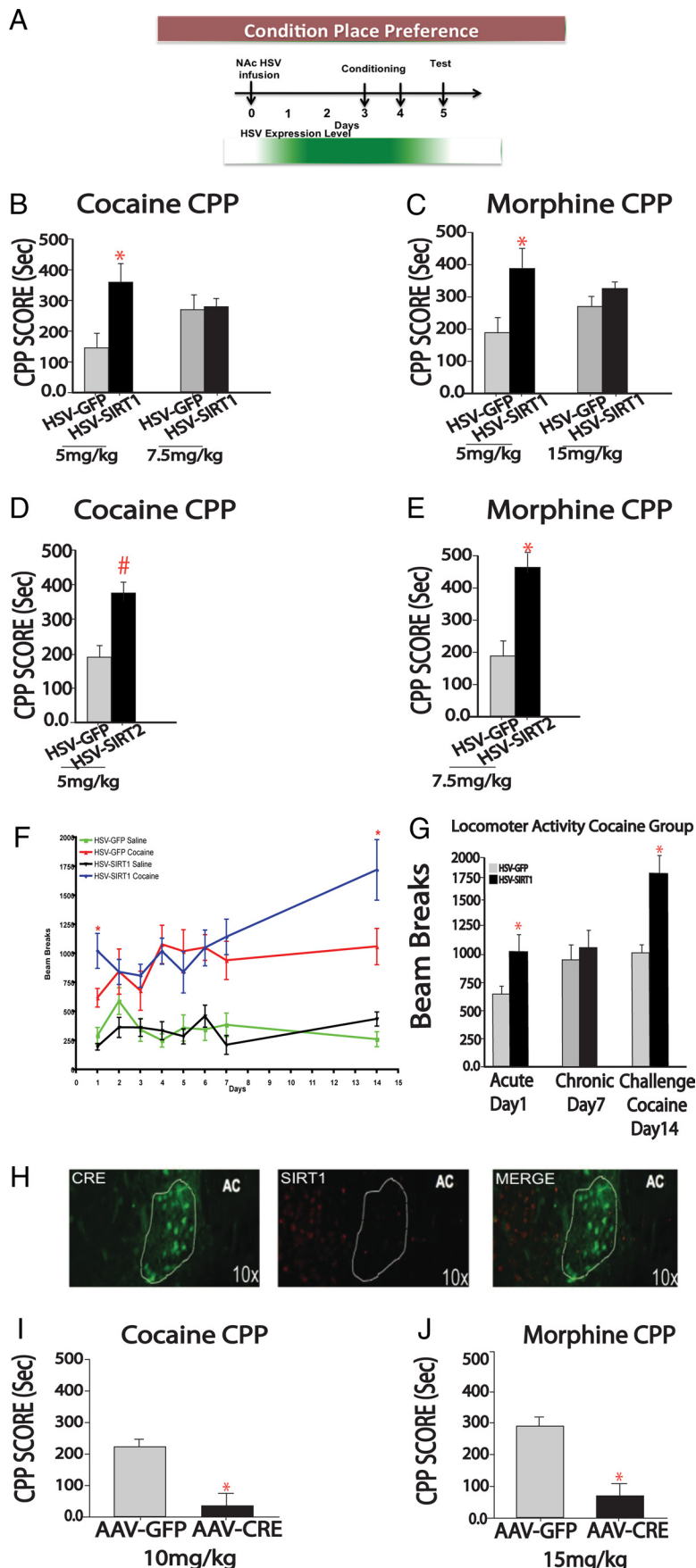


Figure 5. SIRT1 regulates reward-related behaviors. **A**, Temporal expression profile of HSV vectors (based on semiquantitative measures of GFP and transgene expression) and CPP paradigm. Maximal HSV expression occurs within 1 d postinfusion, continues

chromatin-modifying enzymes, which are also altered in NAc in response to chronic cocaine or morphine exposure, including class I and II HDACs, the histone methyltransferase G9a, and the DNA methyltransferase DNMT3a (Robison and Nestler, 2011; Rogge and Wood, 2013).

Regulatory network controlling SIRT expression

We investigated potential upstream transcriptional mechanisms underlying the induction of SIRT1 and SIRT2 in NAc by cocaine. We focused on the transcription factor ΔFosB based on a previous study that found increased ΔFosB binding to the *Sirt2* gene promoter after chronic exposure to cocaine (Renthal et al., 2009). To follow up these findings, we used qChIP to show that chronic cocaine and chronic morphine administration increase ΔFosB binding to the *Sirt1* and *Sirt2* promoters in NAc. Since cocaine and morphine are known to induce ΔFosB in NAc to comparable degrees (Nestler, 2008), it is not surprising that both drugs induce similar increases in ΔFosB to the two promoters. However, the fact that chronic morphine administration does not also induce *Sirt2*, despite the increase in ΔFosB binding, suggests that morphine, but not cocaine, induces other chromatin changes at the *Sirt2* gene, which then oppose ΔFosB’s action, a possibility that now requires investigation. To directly determine whether

through day 4, and dissipates thereafter, consistent with previous studies (Barrot et al., 2002). **B, C**, Cocaine and morphine CPP was studied after intra-NAc injections of HSV-GFP or HSV-SIRT1. Mice expressing HSV-SIRT1 spent more time in the cocaine-paired chamber relative to mice expressing HSV-GFP ($t = 2.267$, $df = 13$) at a lower but not a higher dose of cocaine. Similar results were seen for morphine ($t = 2.276$, $df = 14$). Cocaine and morphine CPPs in animals overexpressing HSV-SIRT2. **D**, For cocaine (5 mg/kg) CPP, mice expressing HSV-SIRT2 tended to spend more time in the cocaine-paired chamber relative to mice expressing HSV-GFP. **E**, For morphine (7.5 mg/kg) CPP, mice expressing HSV-SIRT2 spent more time in the morphine-paired chamber relative to mice expressing HSV-GFP ($t = 2.578$, $df = 13$). The 7.5 mg/kg dose was chosen as intermediate between the two doses used for the experiment in **E, F, G**, HSV-SIRT1 enhances locomotor sensitization to cocaine (7.5 mg/kg). Acute cocaine administration increases locomotor activity in animals expressing HSV-SIRT1 relative to their GFP controls ($n = 7–9$). Cocaine challenge (7.5 mg/kg) 7 d after the last chronic cocaine injection increases locomotor activity in animals expressing HSV-SIRT1 relative to their GFP controls ($t = 2.234$, $df = 13$). **H**, Immunohistochemistry shows an ablation of SIRT1 expression in NAc of floxed *Sirt1* mice injected intra-NAc with HSV-Cre. **I, J**, Animals injected with AAV-Cre exhibited decreased CPP for cocaine administration (10 mg/kg; $t = 2.399$, $df = 15$) and for morphine administration (15 mg/kg; $t = 2.302$, $df = 13$). * $p < 0.05$, # $p < 0.09$.

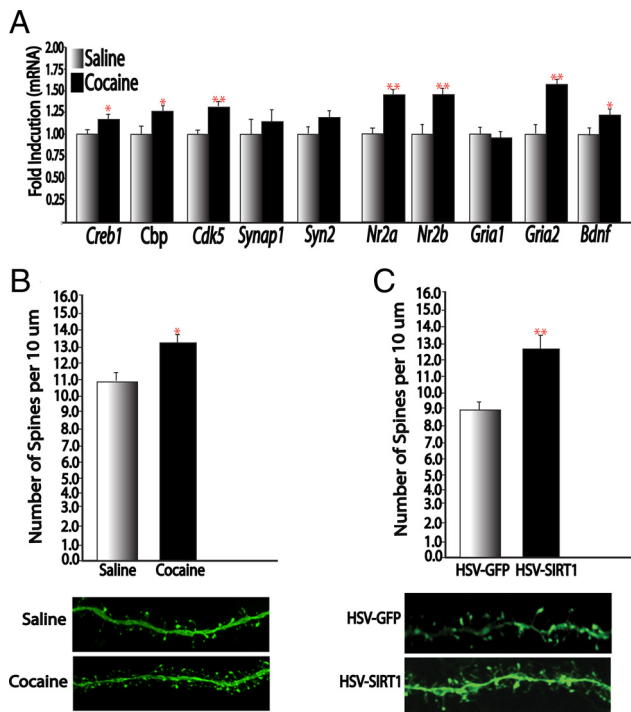


Figure 6. SIRT1 regulates dendritic spine density and expression of plasticity-related genes in NAc. **A**, qPCR analysis of NAc of mice treated for 7 d (20 mg/kg) with cocaine or saline for plasticity-related genes ($n = 7-8$). **B**, **C**, Cocaine increases dendritic spines on NAc medium spiny neurons ($n = 11$ dendrites in 4–5 animals; **B**), an effect mimicked by HSV-SIRT1 ($n = 25-28$ dendrites in 4–5 animals; **C**). * $p < 0.05$, ** $p < 0.01$.

Δ FosB mediates SIRT1 and SIRT2 induction seen after cocaine exposure, we overexpressed Δ FosB in NAc using AAV vectors, and observed increases in both SIRT1 and SIRT2 expression levels. In contrast, overexpressing Δ JunD, a dominant-negative antagonist of Δ FosB, blocked the induction of SIRT1 by cocaine. Together, these data implicate Δ FosB as an essential upstream transcriptional regulator of the sirtuin signaling pathway.

Given the diversity of signaling pathways that regulate sirtuins in other experimental systems, we explored whether some of these other mechanisms occur in our drug abuse models. For example, we were interested in sirtuin regulation at the post-transcriptional level, which has gained wide attention as a critical control point in regulating protein availability (Cheadle et al., 2005). Thus, the ubiquitously expressed RNA binding protein HuR contains three RNA recognition motifs that can bind to the 3' UTR of the *Sirt1* transcript (Abdelmohsen et al., 2007), which increases its half-life in cultured cells, whereas knockdown of HuR has the opposite effect (Abdelmohsen et al., 2007). We found increased *HuR* mRNA levels in NAc after chronic cocaine exposure by qPCR on independent tissue samples, which were shown at the protein level after chronic cocaine or morphine administration via Western blotting. We also explored the possibility that cocaine might regulate SIRT1 post-translationally by analyzing the expression of SUMO-1, which is reported in other systems to enhance the catalytic activity of SIRT1 (Yang et al., 2007). We discovered that chronic cocaine exposure increases SUMO-1 expression in NAc at both the mRNA and protein levels, an effect shown for chronic morphine exposure as well. These findings support the speculation that, in addition to the Δ FosB-mediated increase in *Sirt1* transcription after chronic drug exposure, the increase in HuR and SUMO-1 might, respectively, stabilize the *Sirt1* transcript and

increase SIRT1 deacetylase activity. While further work is required to directly study these possibilities, our results—by demonstrating robust cocaine and morphine regulation of HuR and SUMO-1 expression in NAc—implicate these proteins in drug action for the first time.

Sirtuins regulate behavioral responses to cocaine and morphine

To directly test the influence of SIRT1 and SIRT2 induction in the NAc on behavioral responses to the drugs, we overexpressed either sirtuin in this brain region with HSV vectors or, conversely, we reduced SIRT1 levels locally in NAc using AAV-Cre in floxed *Sirt1* mice, and assessed cocaine and morphine CPPs. We found that overexpression of SIRT1 or SIRT2 in NAc enhances both cocaine and morphine CPPs, whereas local knockdown of SIRT1 in NAc attenuates cocaine and morphine reward. We also found that SIRT1 overexpression in NAc enhances locomotor responses to cocaine. These observations are consistent with our earlier report that systemic administration of a nonselective activator of sirtuins promotes behavioral responses to cocaine (Renthal et al., 2009), and specifically implicates regulation of SIRT1 and SIRT2, acting in NAc per se, in mediating these effects. These behavioral actions of SIRT1 and SIRT2 are consistent with extensive work showing that Δ FosB induction in NAc likewise enhances behavioral responses to drugs of abuse (Nestler, 2008).

SIRT1 regulates dendritic spine density and expression of plasticity-related genes in NAc

A prominent neural adaptation linked to enhanced behavioral responses to cocaine is an increase in the density of dendritic spines on NAc medium spiny neurons (Robinson and Kolb, 2004; Kalivas, 2009; Russo et al., 2010). Further strengthening our hypothesis that SIRT1 is an essential regulator of cocaine-mediated plasticity, we found that overexpression of SIRT1 is sufficient to induce an increase in dendritic spines on NAc neurons. This increase in dendritic spine density in NAc when SIRT1 is overexpressed complements the observation that knockout of SIRT1 decreases dendritic arborizations and spine density in hippocampus (Michán et al., 2010). In concert with the induction of NAc dendritic spine density by SIRT1 is the induction of numerous plasticity-related genes that have been associated with cocaine regulation of synaptic function. Further work is now needed to directly study the involvement of these various genes in mediating the behavioral and structural adaptations of SIRT1. However, it should be noted that, while most studies correlate increased NAc spine density with sensitized behavioral responses to cocaine (Robinson and Kolb, 2004; Russo et al., 2009; Maze et al., 2010; Dietz et al., 2012; Robison et al., 2013), several studies have found opposite effects (Russo et al., 2010), which highlights the need for additional research in this area. Another major paradox in the field is the observation that chronic administration of morphine or other opiates decreases the density of dendritic spines on NAc medium spiny neurons despite the fact that these drugs, like cocaine, produce enhanced reward and locomotor responses (Robinson and Kolb, 2004; Russo et al., 2010). While current efforts are aimed at understanding this paradox, our data would suggest that mechanisms independent of SIRT1 mediate the structural plasticity to opiates.

Conclusions

The results of the present study demonstrate the concerted induction of sirtuin activity in the NAc in response to chronic cocaine or morphine administration, and establish that such induction in turn feeds back to enhance an animal's behavioral

responses to the drugs. These findings contribute to our understanding of the pathophysiology of addiction, and raise the possibility that the host of pharmacological agents being developed to target sirtuins warrant investigation as novel treatment agents for addiction syndromes.

Notes

Supplemental material for this article is available at <http://neuroscience.mssm.edu/nestler/nidappg/chromatingenedatabase.html>. This represents the ChIP-Seq dataset. This material has not been peer reviewed.

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