Hypocretin/Orexin Neurons Contribute to Hippocampus-Dependent Social Memory and Synaptic Plasticity in Mice

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Hypocretin/orexin (Hcrt)-producing neurons in the lateral hypothalamus project throughout the brain, including to the hippocampus, where Hcrt receptors are widely expressed. Hcrt neurons activate these targets to orchestrate global arousal state, wake-sleep architecture, energy homeostasis, stress adaptation, and reward behaviors. Recently, Hcrt has been implicated in cognitive functions and social interaction. In the present study, we tested the hypothesis that Hcrt neurons are critical to social interaction, particularly social memory, using neurobehavioral assessment and electrophysiological approaches. The validated “two-enclosure homecage test” devices and procedure were used to test sociability, preference for social novelty (social novelty), and recognition memory. A conventional direct contact social test was conducted to corroborate the findings. We found that adult orexin/ataxin-3-transgenic (AT) mice, in which Hcrt neurons degenerate by 3 months of age, displayed normal sociability and social novelty with respect to their wild-type littermates. However, AT mice displayed deficits in long-term social memory. Nasal administration of exogenous Hcrt-1 restored social memory to an extent in AT mice. Hippocampal slices taken from AT mice exhibited decreases in degree of paired-pulse facilitation and magnitude of long-term potentiation, despite displaying normal basal synaptic neurotransmission in the CA1 area compared to wild-type hippocampal slices. AT hippocampal slices had lower levels of phosphorylated cAMP response element-binding protein (pCREB), an activity-dependent transcription factor important for synaptic plasticity and long-term memory storage. Our studies demonstrate that Hcrt neurons play an important role in the consolidation of social recognition memory, at least in part through enhancements of hippocampal synaptic plasticity and cAMP response element-binding protein phosphorylation.

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Behavioral/Cognitive
(CREB), an activity-dependent gene transcription. Neurotransmitters such as dopamine, serotonin, and acetylcholine, which enhance memory, induce CREB phosphorylation (Bourchuladze et al., 1994; Silva et al., 1998; Kogan et al., 2000; Shirayama and Chaki, 2006); however, little is known about the function of Hcrt neurons in social interaction and memory, and their physiological roles in the modulation of hippocampal synaptic plasticity and CREB activity.

In the present study, we studied the role of Hcrt in social interaction, especially social memory, using orexin/ataxin-3-transgenic (AT) mice, in which Hcrt neurons degenerate by 3 months of age (Hara et al., 2001). We investigated three components of social interaction (Crawley, 2004; Riedel et al., 2009): (1) sociability: social approach behavior toward unfamiliar conspecifics; (2) social novelty: the preference to interact with a novel conspecific compared with a familiarized one; and (3) social memory: the retention of social recognition over a period of time between sociability and social novelty tests. We found that, compared with their wild-type (WT) littermates, AT mice exhibited normal sociability and social novelty, but had shorter social recognition memory. Attenuated paired pulse facilitation (PPF) and long-term potentiation (LTP) and decreased phosphorylated CREB (pCREB) levels in the AT hippocampus compared with the WT hippocampus may underlie mechanisms of the long-term social memory deficit in AT mice.

Materials and Methods

Animals. Adult male and female mice (AT and WT C57BL/6 littermates, 3–6 months of age) were used. Animals were housed group in standard laboratory conditions and kept on a normal light/dark cycle (lights on at 7:00 A.M.; lights off at 7:00 P.M.). All studies were conducted in accordance with protocols approved by the Institutional Animal Care and Use Committees at AfaSci and SRI International, and the NIH Guide for the Care and Use of Laboratory Animals. Production of transgenic mice followed the procedure described previously (Hara et al., 2001). Presence and copy numbers of the transgene in the offspring were identified by PCR analysis of tail DNA.

Immunohistochemistry and home cage activity monitoring. Hcrt-positive neurons were detected with goat anti-orexin-B antibody (1:100; Santa Cruz Biotechnology) and using the optical fractionator method on epifluorescence photomicrographs (Axiovert inverted fluorescent microscope; Zeiss). The SmartCage system (AfaSci) was used for automated analysis of spontaneous activity in the animal's homecages (L 29.8 cm, W 18.0 cm, H 12.8 cm; Allentown) as described previously (Khroyan et al., 2012). The home cage activity variables (locomotion, travel distance, velocity, and rear-ups) were determined by photobeam breaks and were analyzed automatically using CageScore software (AfaSci). Both AT and WT mice were simultaneously assessed for 48 h during the light/dark cycle and the second 24 h record was plotted in hourly blocks and presented in Figure 1.

Automated two-enclosure home cage social test. The two-enclosure home cage tests (Khroyan et al., 2012) were mostly conducted during the light phase, and several experiments were conducted during the first 3 h of the dark phase as specified in the figure legends. Two transparent Plexiglas enclosures (W 8 × D 6 × H 12 cm) with a metal mesh floor were attached at both end walls of a fresh home cage. The social interaction test consisted of three 10 min sessions based on published procedures (Crawley, 2004; Nadler et al., 2004): (1) habituation, in which the test subject freely explored a home cage with two empty enclosures (nonsocial cues); (2) sociability test, in which the test subject investigated a stimulus mouse designated “Stranger 1,” taken from a home cage different from the test subject’s one. The Stranger 1 was pseudorandomly placed in one of the enclosures; and (3) social novelty test, in which the test subject had the choice of what to investigate, Stranger 1 or a novel mouse designated “Stranger 2,” which was taken from another different home cage, and placed into the empty enclosures. To measure social memory, a delay of 1 min, 10 min, 1 h, 2 h, or 24 h between the end of sociability and beginning of social novelty sessions was imposed. During the delay period, the test subject and Stranger 1 were separated and returned to their own homecages (Nadler et al., 2004; Riedel et al., 2009). The measures of total occupancy time, active time, and traveling distance in each virtual zone were automatically quantified by the SmartCage (Khroyan et al., 2012). The enclosures occupied areas of the homecage that were designated as zones 1 and 3 (zones 1 and 3 were of equal size, whereas the middle zone, zone 2, was twice the size, representing the neutral open area between the two enclosures). All three parameters reflect the same social interaction patterns, but occupancy time is the most sensitive and robust quantification, consistent with the conventional “three-chamber” apparatus test (Crawley, 2004; Nadler et al., 2004).

Nasal administration of Hcrt-1 to mice. The AT and WT mice were nasally dosed with Hcrt-1 (Orexin-A) following the Frey method of nasal administration in rodents (Hanson and Frey, 2008). Mice were individually placed in a supine position while under isoflurane anesthesia within a nose cone. A roll of gauze pad was inserted under the dorsal neck to extend the head back toward the supporting surface. The upper surface of the neck was kept horizontal throughout the dosing procedure to maintain the drug solution in the nasal cavity and to minimize dripping down the nasopharynx. A 2 μl drop of either Hcrt-1 or deionized water was delivered off of the tip of a small pipette and presented to one side of the nares while occluding the opposite naris. One drop of solution was administered every 2 min, alternating between each naris. After each nasal application, the mouse was returned to the nose cone to ensure maintenance of anesthesia. Each naris received 2 drops of solution, totaling a combined volume of 8 μl of Hcrt-1 (0.8 nmol/mouse, n = 14 AT and n = 14 WT) or deionized water (n = 13 AT, n = 14 WT). After completion of the administration, the mice were removed from anesthesia and allowed to regain full consciousness in a fresh home cage. Thirty minutes after each nasal application, each mouse was subjected to the social memory test with a 10 or 120 min delay between sociability and social novelty testing sessions, as described above.

Direct contact social behavioral test. Procedures described previously (Thor and Holloway, 1986; Kogan et al., 2000) were adapted for this study. Individual test subjects were placed into fresh mouse homecages in an observation room under dim light and allowed to habituate to the new environment for 15 min. A stimulus mouse (same sex and similar age as the test subject) taken from a different home cage was introduced into the cage with the test subject as a social cue for an initial interaction trial of 5 min. The intervals between the initial trial and social memory test were 30 min, 3 h, or 24 h as specified in the figure legends. After the selected separation period, the test subject was returned to the test cage with either the familiar or a novel stimulus mouse for the 5 min recognition test. The experimenters manually scored the time that the test subject spent investigating the social cue. Investigation behaviors include contact with any part of the stimulus mouse’s body surface by sniffing, grooming, licking, or pawing and close following (within 1 cm) according to published criteria (Thor and Holloway, 1986; Kogan et al., 2000).

Olfactory habituation/dishabituation test. The ability to detect odors and discriminate between odors was assessed as described previously (Yang and Crawley, 2009). The test mouse was placed into a fresh home cage with a clean metal rack. A clean, dry cotton-tipped wooden applicator (Medline) was inserted through the water bottle hole of the metal rack and fixed in place. Olfactory testing for nonsocial odors was in the following sequence: (1) distilled water, (2) almond extract fresh solution (1:100 dilution; Vanns Spices), and (3) banana extract fresh solution (1:100 dilution; Vanns Spices). Olfactory testing for social odors was done by wiping the applicator in a mouse cage unfamiliar to the test mouse. Each odor was exposed three times consecutively. The experimenters manually scored the cumulative time spent sniffing the tip during the 2 min trial.

Acoustic startle and prepulse inhibition experiments. Acoustic startle and prepulse inhibition were conducted using the SR-Lab startle response system (San Diego Instruments) following a procedure described previously (Geyer and Dulawa, 2003). Mice were placed in a Plexiglas cylinder resting on the sensor platform within a sound-attenuated and ventilated chamber. Acoustic startle stimuli, prepulse stimuli, and continuous background white noise of 65 dB were delivered via a high-frequency speaker. After an acclimation period of 5 min in the containment cylinder, mice were exposed to startle stimuli at five inten-
Data were collected using DigiData 1332 and analyzed with fEPSP evoked by stimulation of the Schaffer collateral and commissural fibers. Data were recorded using an AxoClamp2B amplifier and pClamp 10.3 software (Molecular Devices). Field EPSP (fEPSP) in Schaffer collateral-CA1 synapses was recorded using a glass microelectrode filled with the ACSF. Biphasic current pulses (0.2 ms) were delivered through a concentric bipolar stimulator (FHC). Both recording and stimulus electrodes were placed in parallel on stratum radiatum separated by ~500 μm to record fEPSP evoked by stimulation of the Schaffer collateral and commissural fibers. Data were collected using DigiData 1332 and analyzed with Clampfit (Molecular Devices). Input-output (I-O) curves were obtained using stimulus intensity from the threshold to a strength evoking the maximum response. The fEPSP slope was measured from the initial phase of the negative wave. The I-O curve was constructed by plotting the mean fEPSP slope against the fiber voltage amplitude. The fiber voltage amplitude was binned at 0.2 mV across slices for clarity of presentation. For the basal synaptic activity, test pulse intensities were adjusted to 30–35% of the maximal stimulation intensity and delivered at 10 s intervals. The PPF was induced by a pair of stimulations at intervals of 20, 50, or 100 ms. High-frequency stimulation (HFS) consisted of 4 trains of 100 pulses (0.2 ms pulse duration, 100 Hz) with an intertrain interval of 20 s. LTP was plotted as the percentage changes in the fEPSP slope after HFS relative to the basal slope of 5 averaged consecutive fEPSP (mean ± SEM) just before HFS.

Measurement of CREB and pCREB. Proteins were extracted from fresh hippocampi taken from the AT and WT littermates after the social interaction tests. Total CREB and pCREB were detected using antibodies #9197S and #9198S, respectively (Cell Signaling Technology) as described previously (Tardito et al., 2009). β-actin was obtained from Sigma. In each lane, 30 μg of protein was subjected to SDS-PAGE using 4–15% Ready Gel (Bio-Rad) for 1.5 h. Protein bands were transferred from the gel to PVDF membranes (Millipore) for 1 h. After the membrane was blocked with 5% nonfat dry milk (Bio-Rad) in PBS/0.05% Tween 20, primary antibodies were incubated overnight at 4°C, followed by exposure to horseradish peroxidase-conjugated secondary antibody (1: 2000; Cell Signaling Technology) for 1 h. The membrane was then incubated with anti-β-actin antibodies (1:10,000; Sigma) as an even-loading control of total protein. Membranes were scanned using the Typhoon trio (GE Healthcare). Optical densities of CREB and pCREB were analyzed from Western blots with normalization to β-actin.

Data analysis. The experiments were performed with genotypes and treatments blinded to the experimenters. All data are presented as mean ± SEM and were subjected to statistical analysis using within genotype repeated-measures ANOVA, with the factor of type as the between-genotype variable and time interval as the repeated measure. For two groups, paired or unpaired Student’s t tests with two tails. For multiple groups, post hoc comparisons were done using Bonferroni’s correction. 

Results

AT mice display normal sociability in the two-enclosure homoeage test

Compared with their WT littermates, AT transgenic mice had substantial Hcrt-positive cell loss (Fig. 1a,b) and exhibited significantly lower spontaneous activity as indicated by active time (Fig. 1c; ANOVA, F_{1,14} = 0.040, Fisher’s PLSD at recording hour 1, p = 0.048, and at hour 8, p = 0.015), and distance traveled (Fig. 1d, hour 1, p = 0.025, and hour 8, p = 0.014). However, AT mice did not show significant differences in rearing (Fig. 1e) or traveling speed (Fig. 1f) in the dark phase measured using the SmartCage (Fig. 1c–f), which is consistent with the previously reported phenotype (Hara et al., 2001). We used the validated

Testing for two tails. For two groups, paired or unpaired Student’s t tests were used, as appropriate. Multiple-group comparisons were analyzed using two-way repeated measures ANOVA with drug or genotype as the between-genotype variable and time interval as the repeated measure. If the ANOVA was statistically significant, Fisher’s PLSD was used to determine group differences. Data from the two components of the social behavior test (sociability and social novelty) were analyzed using within genotype repeated-measures ANOVA, with the factor of cage side (e.g., zone 1 vs 3). The difference was considered significant when p < 0.05.
two-enclosure homecage testing procedure and quantified mouse behavior using the automated SmartCage system (Khroyan et al., 2012). Under these conditions, the test subject freely explored the fresh homecage with nonsocial cues (i.e., identical empty enclosures) or social cues (i.e., stranger mice that were taken from different homecages). During habituation, most test mice (AT and WT) spent an equal amount of time in zones 1 and 3, indicating an equal amount of interest or no bias toward either nonsocial cue. During the sociability session, all test subjects spent more time investigating Stranger 1 compared with the nonsocial cue (i.e., empty enclosure, Fig. 2a,b). A paired Student’s t test with genotype: WT: \( t_{(7)} = 4.47, p = 0.003 \); AT: \( t_{(7)} = 5.59, p = 0.001 \). During the social novelty test without a delay between the completion of sociability and the initiation of social novelty test, all test subjects spent more time investigating Stranger 2 than the familiarized Stranger 1 (Fig. 2b). AT mice displayed preferential Stranger 1 in a social memory test with a 10 min delay (Fig. 2c). Social memory was equivalent between the genotypes. However, AT mice retained short-term social memory after a 1 min delay, whereas WT mice preserved their memory after a 10 min delay, indicating that there is a significant delay in the social memory observed in AT mice could be due to a reduction in arousal or alertness levels during the circadian light phase. Separate cohorts of AT and WT littermates were tested for their social interaction and memory in the first 3 h of the dark phase using the same 2-enclosure homecage test. As in the light phase, AT mice and their WT littermates exhibited normal sociability and social novelty when there was no delay (Fig. 3a,b). However, AT mice exhibited deficits in long-term social memory after a 1 min delay, whereas WT mice retained short-term social memory after a 10 min delay, indicating that WT mice did not exhibit a loss of social memory after a 10 min delay, whereas WT mice preserved their memory after a 10 min delay (Fig. 3c). AT mice still exhibited a loss of social memory after a 10 min delay, whereas WT mice preserved their memory after a 10 min delay (Fig. 3c) and 1 h delay (data not shown) as tested in the light phase. WT mice failed to show preference for novel social cues after a delay of 2 h (Fig. 3e), indicating that WT mice did not dramatically improve social memory tested with this delay time point during the dark phase.

Nasal administration of Hcrt-1 improves social memory in AT mice
To test our hypothesis that a supplement of exogenous Hcrt in the brain of AT mice can lead to an overall improvement in social memory, AT mice were treated with Hcrt-1 via nasal administration, a method that has been shown to deliver a variety of peptides effectively into the brains of rodents (Hanson and Frey, 2008). Thirty minutes after nasal administration of either vehicle, deionized water (\( n = 13 \)) or Hcrt-1 (0.8 nmol, \( n = 14 \)), the AT mice were subjected to the sociability test followed by a 10 min delay social memory test. The AT mice that were treated with vehicle spent significantly more time investigating Stranger 1 than the empty enclosure, indicating normal sociability. However, these mice did not shift their investigation to Stranger 2 over the familiarized Stranger 1 in a social memory test with a 10 min delay (Fig. 4a). AT mice treated with Hcrt-1 displayed not only a bias toward Stranger 1 (Fig. 4b). This result indicates that there are no significant differences in sociability and social novelty between the genotypes.
normal sociability, but also a distinct shift in their investigation to Stranger 2 from familiarized Stranger 1 (Fig. 4c-f). Interestingly, Hcrt-1-treated WT mice did not produce a significant increase in social memory as tested with a 120 min delay compared with nasal vehicle treatment (data not shown) or nonnasal treatment (Fig. 2e).

**AT mice display normal social approach behavior but social memory deficit in direct contact social test**

To confirm the normal sociability of AT mice, we performed a direct contact social test with manual scoring of the test subject’s interaction with a stimulus mouse taken from a different home cage as a social cue. The onset and duration of social interaction was indistinguishable between AT and WT littermates in the initial trial, confirming that both genotypes have normal sociability (61.0 ± 4.5 s in AT vs 62.1 ± 6.7 s in WT, n = 8/group, unpaired Student’s t test, t(12) = 14, p > 0.05; Fig. 5a,b).

To verify deficits in the social memory of AT mice, the test subjects were exposed repeatedly to the same initial social cue with varying separation periods. Previous studies have suggested that the test subject’s investigation time in the initial trial reflects sociability, whereas a reduction in investigation time during the second exposure to the same social cue is correlated with social recognition retention (Thor and Holloway, 1986; Kogan et al., 2000). In the initial exposure, the test subjects spent ~62.1 ± 6.7 s investigating the social cue, as described above (Fig. 5a). In the second exposure to the same social cue, after a separation period of 24 h, 3 h, or 30 min, the WT test subjects decreased their investigation time toward the same social cue as the separation time reduced. The WT mice could at least recognize and remember the familiarized social cue after separation of up to 3 h (and not beyond 24 h), as indicated by a significant decrease in investigation time compared with the initial trial (Fig. 5c). In contrast, the AT test subjects displayed decreased investigation time toward the familiarized social cue when the separation period was 30 min (t(30) = 6.17, p < 0.001, and at 3 h t(30) = 2.98, p < 0.05). In contrast, the AT test subjects displayed decreased investigation time toward the familiarized social cue when the separation period was 30 min (t(30) = 6.16, p < 0.001), but the AT mice did not reduce their investigation time after a 3 h delay (t(27) = −1.25, p > 0.05; Fig. 5b), which confirms that they had a shorter recognition retention time compared with WT mice.

To eliminate the possibility that decreased investigation time in repeated trials was due to a decrease in interest/motivation of social interaction or locomotor activity, which could confound social memory readout, we tested social recognition retention with different stimulus mice between the initial trial and the second test. As shown in Figure 5c, both genotypes showed investi-
genotype). Paired Student’s t test, with the exception of the first banana exposure (unpaired Student’s t test, \( t_{(14)} = -2.12, p < 0.05 \)), all data points were not significantly different between genotypes. Furthermore, the olfactory habituation/dishabituation patterns were almost identical to the published data (Yang and Crawley, 2009). Therefore, we conclude that deficits in social memory in AT mice were not caused by differences in olfactory discrimination ability (Fig. 6a).

Alternatively, auditory ability differences between AT and WT mice could also influence social interaction. To eliminate this possibility, we assessed acoustic startle responses evoked by a series of sound intensities (80–120 dB) with an increment of 10 dB. There were no significant differences in startle responses (Fig. 6b) or in prepulse inhibition between the two genotypes under our experimental conditions (Fig. 6c).

**Figure 5.** AT mice showed normal social interaction but with a social memory deficit compared with WT littermates in a direct contact social test. a, Test WT mice displayed social approach behavior toward the stimulus mouse (sociability) and reduced the approach behavior toward the reexposed social cue mouse 30 min and 3 h after initial contact as measured by a reduction in investigation time (\( n = 8 \) per genotype). Paired Student’s t test: *\( p < 0.05 \); **\( p < 0.01 \). After a 24 h delay after the initial interaction, the test mouse spent the same amount of time investigating the same social cue mouse as in the initial trial. b, Test AT mice displayed social approach behavior toward the stimulus mouse (sociability) and reduced investigation time toward the same social cue mouse 30 min after the initial interaction (\( n = 8 \) per genotype). Paired Student’s t test: *\( p < 0.05 \); **\( p < 0.01 \). After a delay of 3 or 24 h after the initial interaction, the test mouse spent a similar amount of time investigating the reexposed social cue as in the initial interaction. c, AT and WT mice show similar investigation times toward two different, unfamiliar social cue mice between the initial trial and a second test after a 3 h delay. In the initial contact, the test subject displayed social approach behavior, as indicated by time investigating the stranger mouse. Three hours later, the same test subject was tested for its sociability toward a novel mouse (taken from another home cage). Subjects were 3- to 4-month-old male mice (\( n = 8 \) per genotype). Paired Student’s t test, \( p > 0.05 \).

**Figure 6.** Comparison of olfactory and auditory reflex functions in AT mice and their WT littermates. a, Olfactory habituation/dishabituation ability was measured as time spent sniffing a sequence of novel nonsocial (i.e., water, almond, and banana) and social odors (i.e., unfamiliar mouse cage odor) contained in cotton swab tips. Except for banana 1 (unpaired Student’s t test, *\( p < 0.05 \)), all data points were not significantly different between genotypes. b, Startle responses evoked by different sound intensities (80–120 dB). There were no significant differences between the genotypes (\( n = 8 \) per group). c, Prepulse inhibition induced by a constant startle stimulus (120 dB, 40 ms), which was preceded by a prepulse (70–90 dB, 20 ms) with an interval (100 ms) between the prepulse and startle stimuli. There were no significant differences between the genotypes. Subjects were 3- to 4-month-old male mice (\( n = 8 \) per genotype).

**AT mice display deficits in hippocampal synaptic plasticity**

Because social recognition memory is dependent on hippocampal long-term synaptic plasticity (Kogan et al., 2000), we studied hippocampal synaptic basal activity and plasticity in the CA1 area using extracellular recording techniques as described previously (Xie and Smart, 1994). We first investigated basal synaptic transmission by analyzing the I-O relationships of fEPSPs evoked by various stimulus intensities. There were no significant differences in the I-O curves between AT and WT mice, indicating that AT mice have normal basal synaptic transmission and synaptic coupling efficiency in the Schaffer collateral-CA1 synapses (Fig. 7a).

We next analyzed PPF, a presynaptic form of short-term synaptic plasticity, using intervals of 20, 50, and 100 ms between a pair of stimulation pulses. All three paired pulses of different
shows that with a 20 ms stimulus interval, the slope of the second fEPSP indicates that the two genotypes had a similar level of basal synaptic neurotransmission but a decreased PPF ratio and attenuated LTP in the CA1 area of hippocampal slices compared with WT littermates. Therefore, the smaller magnitude of LTP in AT mice was not caused by differences in basal synaptic strength.

To investigate the functional consequences of the loss of Hcrt neuronal input on long-term synaptic plasticity, we compared LTP in the CA1 area. The results indicated that the magnitude of the maintenance phase of LTP in WT mice was greater than that in AT mice (Fig. 7c). Analysis of the averaged five consecutive slopes of fEPSP in the CA1 in WT mice revealed an increase to 234.6 ± 23.8% of the baseline level at 45 min after HFS (n = 14 slices from 5 mice). In AT mice, the slope of fEPSP in CA1 was 164.3 ± 12.7% of baseline at the same time point (n = 14 slices from 6 mice; unpaired Student’s t test, t(24) = −2.71, p = 0.02). These results indicate that synaptic plasticity in AT slices was impaired compared with WT mice. Before HFS, the basal level of fEPSP slopes for WT and AT mice was −0.14 ± 0.01 and −0.19 ± 0.02 mV/ms, respectively (t(26) = −1.87, p = 0.07). Therefore, the smaller magnitude of LTP in AT mice was not due to differences in the basal level of fEPSP between genotypes (Fig. 7c).

AT hippocampi have lower levels of pCREB

Lastly, we explored possible signaling pathways related to hippocampal CA1 LTP and social memory in behavior. CREB is a nuclear protein that modulates the transcription of genes with cAMP-responsive elements in their promoters and acts as a universal modulator of processes required for memory formation, especially hippocampus-dependent social memory. Mice deficient in the α and Δ isoforms of the CREB transcription factor (CREBaD2 mice) are impaired in long-term, but not short-term, social memory (Bourtchuladze et al., 1994; Silva et al., 1998; Kogan et al., 2000). We therefore investigated the level of total CREB and pCREB, the active form of CREB in hippocampi, taken from mice after social interaction and memory testing (n = 9 mice per genotype). In both genotypes, CREB immunostaining was found to be distributed heterogeneously in most cell bodies throughout the hippocampus (Fig. 8a). Immunostaining indicated that there were no significant differences in total CREB expression between the genotypes. However, the pCREB level was apparently lower in the hippocampus taken from AT compared with that of WT mice, although most neurons expressed pCREB in both genotypes (Fig. 8a). Optical density analysis on the Western blot from hippocampi from each genotype quantified the CREB and pCREB levels. The quantitative analyses obtained from the Western blots confirmed that there was no significant differences in total CREB expression between the genotypes and revealed that pCREB in the AT hippocampus was significantly lower by 25 ± 7.5% compared with the WT hippocampus (Fig. 8b; n = 9 Western blot bands from nine mice of each genotype, unpaired Student’s t test, t(16) = 2.29, p < 0.05).

Discussion

Social recognition memory, the ability to distinguish and remember familiar from novel conspecifics, is critical for appropriate social behavior. Rodents can retain recognition of recently encountered conspecífics for a certain period of time. In the present study, using AT mice in which Hcrt neurons are genetically ablated (Hara et al., 2001), we obtained the novel result that the endogenous Hcrt system is essential for the formation of long-term social memory. A recent microdialysis study of Hcrt in human brain revealed a dramatic increase in Hcrt levels associated with social interaction (Blouin et al., 2013). Our study supports this clinical observation and defines the physiological role of the
Hcrt system in social memory, but not necessarily in the initiation of social approach, because AT mice displayed normal sociability. Our results suggest that the Hcrt surge upon interaction with conspecifics consolidates storage of information, but may not be critical in the stages of encoding (e.g., processing information) or retrieval (e.g., recalling) of stored information, because AT mice could process and recognize the familiarized social cue and displayed preference to the novel social cue when there was no delay or a short delay (e.g., 1 min) in the memory test.

To assess social interaction in humans and animal models, several behavioral tests have been developed (Kaidanovich-Beilin et al., 2011). The three-chamber apparatus and testing procedure have been used commonly to assess rodents’ sociability, social novelty, and social memory (Crawley, 2004; Moy et al., 2004; Deng et al., 2007). AT mice display decreased active time and locomotion, which could confound social behavior testing, especially when using a large, unfamiliar chamber with physical dividers. We therefore developed and validated a two-enclosure test in a mouse homecage (Khroyan et al., 2012), and showed the social interaction pattern to be comparable to the results obtained previously using the three-chamber test (Crawley, 2004; Moy et al., 2004; Deng et al., 2007). Using this objective testing method, we found that AT mice displayed normal sociability and social novelty, but exhibited a long-term memory deficit.

The social memory of AT mice was shorter than 1 min when the test was conducted in the light phase, but at least 1 min and shorter than 10 min when the task was performed in the dark phase. The improvement in social recognition task performance during the dark phase not only supports the finding of the specific role of Hcrt neurons in reward behavior during the diurnal phase (McGregor et al., 2011), but also suggests that the arousal-promoting role of Hcrt may be crucial in the consolidation of memory storage. Interestingly, the social memory of the WT mice was not enhanced at the time point tested in the dark phase, suggesting a possible “saturated effect” of Hcrt levels in WT mice. Alternatively, enhancement of social memory could be observed in refined delay time points.

Hcrt neurons also produce other neurotransmitters, such as dynorphin, pentraxin, and glutamate, which may influence memory (Chou et al., 2001; Reti et al., 2002; Rosin et al., 2003). Therefore, it is important to determine whether a supplement of exogenous Hcrt in the brain can restore long-term social memory in AT mice. Indeed, we found that nasally administered Hcrt-1 extended social memory in AT mice from <1 min to at least 10 min when tested in the light phase. In contrast, Hcrt-1 did not prolong social recognition retention in WT mice tested with a 120 min delay. These findings are compatible with previous results showing that nasal administration of Hcrt-1 selectively improves sleep-deprivation-caused memory deficits without enhancing memory in normal nonhuman primates (Deadwyler et al., 2007). However, these treatment regimens were acute and whether chronic treatment with Hcrt can enhance memory is worthy of further investigation.

The deficit in long-term social memory in AT mice was corroborated by a conventional direct contact social test. In this test, social recognition memory was inferred by a decrease in the investigation time toward the reexposed social cue (Thor and Holloway, 1986; Kogan et al., 2000). The direct contact allows more engagements between the test subject and the social cue, leading to stronger social stimulation, and therefore longer recognition retention, compared with the two-enclosure test. AT mice exhibited normal sociability toward unfamiliar conspecifics and their social recognition retention lasted at least 30 min but <3 h. In contrast, the WT mice’s recognition retention lasted for at least 3 h and disappeared after 24 h of separation. The AT mouse deficit in long-term memory was not due to any defects in olfactory or auditory ability, because there were no significant differences in the olfactory habituation/dishabituation patterns or acoustic startle responses and prepulse inhibition between the AT and WT littermates.

Social recognition, particularly long-term memory in humans and rodents, relies critically on intact hippocampal structure and function (Corkin et al., 1984; van Wimersma Greidanus and Maigret, 1996; Kogan et al., 2000). The hippocampus receives direct input from Hcrt neurons and expresses Hcrt receptors. Furthermore, Hcrt-mediated activation of noradrenergic and serotonergic systems might also play a role in consolidating memory, because these monoaminergic neurons project abundantly to the hippocampus ( Peyron et al., 1998; Kilduff, 2005; Carter et al., 2009; Eriksson et al., 2010; de Lecea, 2010; Sakurai and Mieda, 2011). We therefore focused on the investigation of hippocampal synaptic plasticity and signaling pathway to explore mechanisms underlying the effects of Hcrt in the storage of social memory.

Although CA1 had normal basal neurotransmission in both AT mice and their WT littermates, both PPF and LTP were
significantly attenuated in the AT hippocampi. These results suggest that short- and long-term synaptic plasticity in the AT hippocampus may be impaired in the absence of Hcrt neuronal input. Conversely, an LTP-like phenomenon in the CA1 area could be induced by the application of exogenous Hcrt-1 in hippocampal slices via activation of several protein kinases, including protein kinase A and C (Selbach et al., 2010). Moreover, Hcrt in the ventral tegmental area has been shown to be critical for the induction of synaptic plasticity that was associated with behavioral sensitization to cocaine in rats (Borgland et al., 2006).

The formation of long-term memory in hippocampus-dependent cognitive tasks, including social recognition, has been found to involve CREB. CREB acts as memory storage and sustains synaptic plasticity in a wide range of synapses, including in the hippocampus. Several neurotransmitters, especially dopamine, serotonin, noradrenaline, and acetylcholine, all of which are known to enhance memory, induce CREB phosphorylation (Bourtchuladze et al., 1994; Silva et al., 1998; Kogan et al., 2000; Thonberg et al., 2002). The Hcrt system has been found to synapse directly and activate the amine systems and to promote noradrenaline and acetylcholine release (Carter et al., 2009; Eriksson et al., 2010; Stanley and Fadel, 2012). Complementary to this indirect “upstream” modulation of CREB, Hcrt can activate CREB directly by acting on Hcrt receptors that are Gq-protein-coupled receptors. This leads to signaling through phospholipase C and calcium-dependent and calcium-independent transduction pathways (Sakurai et al., 1998; Sakurai and Mieda, 2011). Activation of Hcrt receptors also leads to stimulation of protein kinase A and C, as mentioned above (Selbach et al., 2010), and a recent study showed that exogenous Hcrt prolonged CREB phosphorylation via protein kinase C activation (Guo and Feng, 2012). As shown in this study, the level of pCREB in WT hippocampi was significantly higher than in AT hippocampi, whereas there were no significant differences in total CREB between the genotypes.

Social memory underpins and enables appropriate social interaction, which is critical for the establishment of relationships and for the stability of the social networks. Several neuropsychiatric disorders are characterized by disruptions of social recognition or long-term social memory, including schizophrenia, bipolar disorders, depression, attention-deficit hyperactivity disorder, and autism spectrum disorders (McCleure et al., 2003; Riedel et al., 2009; Brotman et al., 2010; Ibi et al., 2010; Kaidanovich-Beilin et al., 2011). Loss of Hcrt neurons causes narcolepsy in humans (Scammell et al., 2001; Siegel et al., 2001). Narcoleptic patients are often anxious about disease-related deficient memory functions (Aguirre et al., 1985; Rogers and Rosenberg, 1990) and modest memory impairments have been described previously (Henry et al., 1993; Naumann et al., 2006). Although narcoleptic patients display normal sociability, whether they have deficits in long-term social memory is currently unknown, and this can be a potential area for clinical investigation.

The present study has uncovered a physiological role of Hcrt neurons in long-term social memory using a transgenic animal model and pharmacological approach. Hcrt neurons apparently are not essential for sociability, but play a critical role in the formation of long-term social memory. This finding is consistent with the notion that the Hcrt system contributes to spatial and nonsocial learning and memory (Jaeger et al., 2002; Akbari et al., 2007; Eriksson et al., 2010; Fadel and Burk, 2010). The mechanisms underlying the effects of Hcrt in the formation of social memory is through, at least in part, enhancing synaptic plasticity and increasing pCREB in the hippocampus. The previously unrecognized role for the Hcrt system in social memory may have clinical implications and our study will foster further integrative research using animal disease models and appropriate tests to identify novel neuronal pathways and therapeutics for the modulation of social interaction and recognition memory.

Notes
Supplemental material for this article is available at http://www.afasci.com/publications/afasci-publications. Automated two-enclosure homecage social test and spontaneous activity and locomotion monitoring using the SmartCage system. This material has not been peer reviewed.

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