

# Altered Sleep Homeostasis after Restraint Stress in 5-HTT Knock-Out Male Mice: A Role for Hypocretins

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Restraint stress produces changes in the sleep pattern that are mainly characterized by a delayed increase in rapid eye movement sleep (REMS) amounts. Because the serotonin (5-HT) and the hypocretin (hcrt) systems that regulate REMS are interconnected, we used mutant mice deficient in the 5-HT transporter (5-HTT<sup>-/-</sup>) to examine the role of 5-HT and hcrt neurotransmissions in the sleep response to stress.

In contrast to wild-type mice, restraint stress did not induce a delayed increase in REMS amounts in 5-HTT<sup>-/-</sup> mice, indicating impaired sleep homeostasis in mutants. However, pharmacological blockade of the hcrt type 1 receptor (hcrt-R1) before restraint stress restored the REMS increase in 5-HTT<sup>-/-</sup> mice. In line with this finding, 5-HTT<sup>-/-</sup> mutants displayed after restraint stress higher long-lasting activation of hypothalamic preprohcrt neurons than wild-type mice and elevated levels of the hcrt-1 peptide and the hcrt-R1 mRNA in the anterior raphe area. Thus, hypocretinergic neurotransmission was enhanced by stress in 5-HTT<sup>-/-</sup> mice. Furthermore, in 5-HTT<sup>-/-</sup> but not wild-type mice, hypothalamic levels of the 5-HT metabolite 5-hydroxyindole acetic acid significantly increased after restraint stress, indicating a marked enhancement of serotonergic neurotransmission in mutants.

Altogether, our data show that increased serotonergic- and in turn hypocretinergic- neurotransmissions exert an inhibitory influence on stress-induced delayed REMS. We propose that the direct interactions between hcrt neurons in the hypothalamus and 5-HT neurons in the anterior raphe nuclei account, at least in part, for the adaptive sleep-wakefulness regulations triggered by acute stress.

## Introduction

Stress or aversive life events are known to alter sleep and wakefulness. Such stress-induced changes in the sleep-wakefulness patterns have been proposed to be part of the process that enables an organism to adapt to emotional inputs (Van Reeth et al., 2000). In rodents, restraint stress (RS) has been found to produce a delayed increase in rapid eye movement sleep (REMS) that may last for up to 10 h (Rampin et al., 1991) and depends on serotonergic mechanisms. Thus, in rats, stimulation of the dorsal raphe nucleus (DR), which increases 5-HT release in the hypothalamus, is sufficient to induce a delayed increase in REMS (Houdouin et al., 1991a,b). Conversely, this phenomenon is abolished in transgenic mouse models with permanent impairment of 5-HT neurotransmission (Boutrel et al., 2002; Adrien et al., 2004; Popa et al., 2008).

Mutant mice that do not express the serotonin transporter (5-HTT<sup>-/-</sup>) exhibit a marked increase in extracellular 5-HT levels (Fabre et al., 2000; Mathews et al., 2004), which, in turn, triggers adaptive changes in 5-HT receptor expression and/or function (for review, see Murphy et al., 2008). In addition, 5-HTT<sup>-/-</sup> mice show exaggerated hypothalamic-pituitary-adrenal (HPA) responses that may correlate with an increased reactivity to stress and anxiety-like behaviors (Hariri and Holmes, 2006). Therefore, 5-HTT<sup>-/-</sup> mutants are a relevant model to explore further the influence of stress on sleep-wakefulness patterns and to elucidate underlying neurobiological mechanisms.

Hypocretins (hcrt)/orexins are two neuroexcitatory peptides, hypocretin-1 (orexin-A) and hypocretin-2 (orexin-B), first isolated from the rat hypothalamus, which derive from a single precursor, the preprohcrt (de Lecea et al., 1998; Sakurai et al., 1998). Both hcrt and 5-HT play essential roles in the control of sleep and wakefulness, notably through an inhibitory influence on REMS generation (Pace-Schott and Hobson, 2002; Lu et al., 2006). Indeed, these peptides and the indolamine exhibit strong interconnections: (1) hypocretin neurons send projections to some raphe nuclei, and, in turn, they receive 5-HT inputs (Peyron et al., 1998; H. S. Lee et al., 2005; Sakurai et al., 2005; Wang et al., 2005; Yoshida et al., 2006), and (2) 5-HT hyperpolarizes hypocretin neurons by activating 5-HT<sub>1A</sub> receptors (Collin et al., 2002; Li et al., 2002; Muraki et al., 2004), whereas hypocretins excite DR 5-HT neurons by acting at hcrt-R1 and hcrt-R2 receptor subtypes

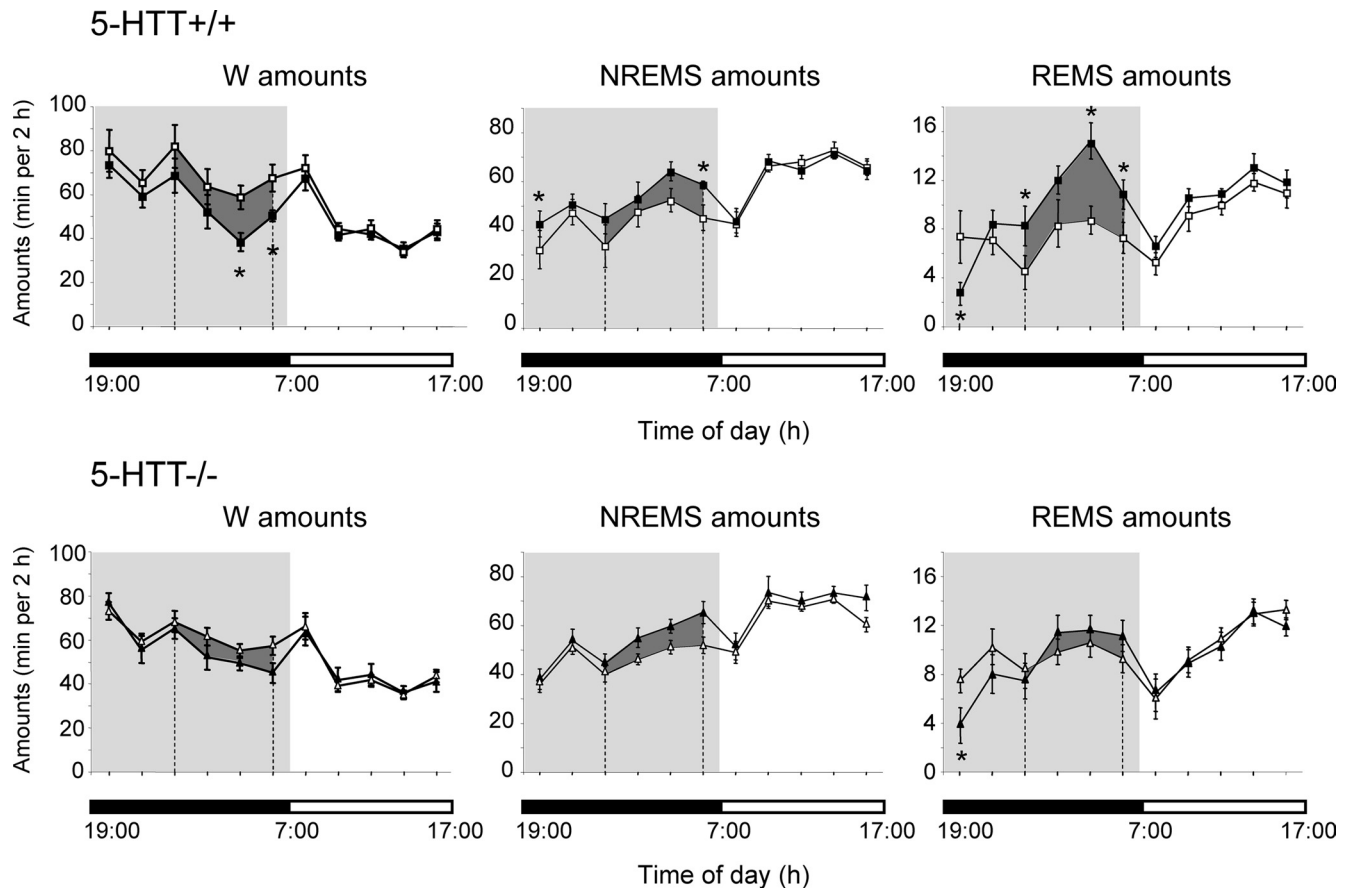
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**Figure 1.** Effects of restraint stress on sleep–wakefulness patterns in *5-HTT<sup>+/+</sup>* and *5-HTT<sup>-/-</sup>* mice. Sleep–wakefulness states across the light/dark cycle under control conditions (open symbols) or after RS (filled symbols) in *5-HTT<sup>+/+</sup>* (squares, top) and *5-HTT<sup>-/-</sup>* (triangles, bottom) mice. Mice were, or were not, restrained for 90 min (5:30 P.M. to 7:00 P.M.) and then monitored for sleep and wakefulness (from 7:00 P.M. on day 1 to 5:00 P.M. on day 2; abscissa). Amounts of vigilance states, expressed as minutes per 2 h, are the mean  $\pm$  SEM of eight animals in each group. The gray area corresponds to the difference between control and RS conditions. \* $p < 0.05$ , significantly different from the control conditions; paired Student's *t* test after ANOVA.

(Brown et al., 2001, 2002; Marcus et al., 2001; Liu et al., 2002; Soffin et al., 2004; Takahashi et al., 2005; Wang et al., 2005; Tao et al., 2006).

On this basis, our aim was to specify the role of 5-HT and hypocretin in stress-altered sleep in an animal model with well characterized alterations in 5-HT neurotransmission, *5-HTT<sup>-/-</sup>* mutants versus paired wild-type (*5-HTT<sup>+/+</sup>*) mice. We notably investigated the physiological role of hypocretin neurons in stress-induced sleep modifications by assessing the effects of hcrt-R1 blockade during stress on subsequent sleep in *5-HTT<sup>-/-</sup>* versus *5-HTT<sup>+/+</sup>* mice. Finally, we also examined the impact of 5-HTT deficiency on serotonergic and hypocretinergic neurotransmissions under control conditions and after RS in both mutants and wild-type mice.

## Materials and Methods

### Animals

Founder 5-HTT knock-out mice were obtained originally from the colony of author K.P.L. (Bengel et al., 1998) and were successively backcrossed (F7) with the Swiss albino CD-1 strain (Charles River). Males of each genotype were born from either heterozygous ( $n = 82$ ) or homozygous ( $n = 114$ ) breeding. Genotyping was performed by PCR (Bengel et al., 1998). Animals were housed in a temperature-controlled room ( $23 \pm 1^\circ\text{C}$ ) under 12 h light/dark cycle (lights on at 7:00 A.M.), with food and water available *ad libitum*. Animal care and experiments were conducted in accordance with the institutional guidelines, which are in compliance with national and international laws and policies (council directive 87-848, October 19, 1987, Ministère de l'Agriculture et de la Forêt, Service

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### Experimental design

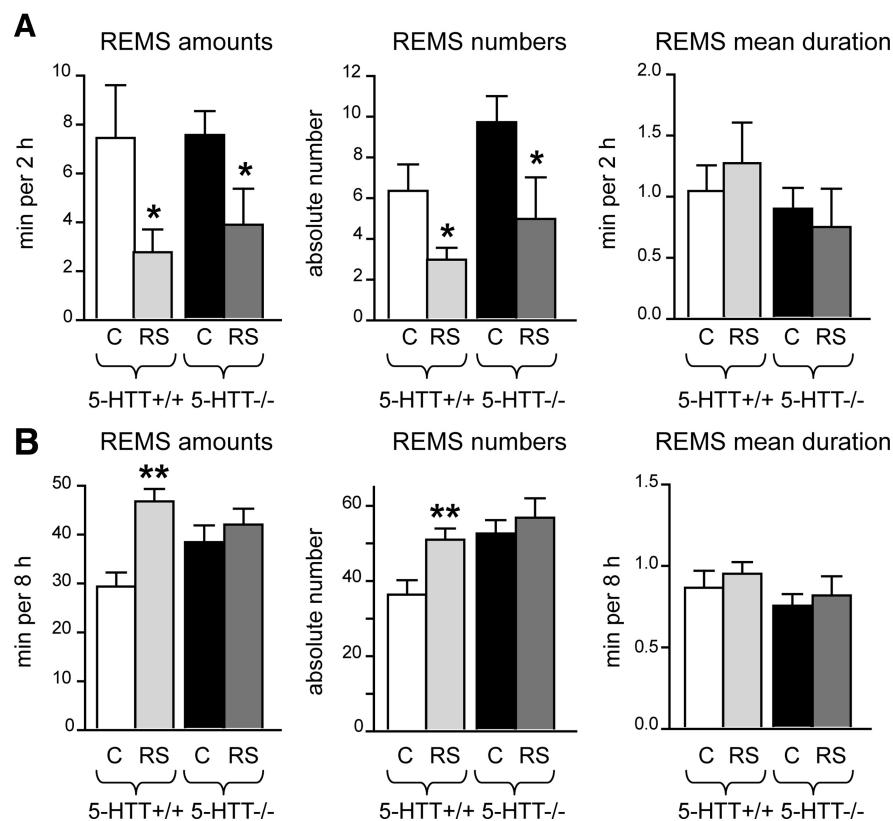
To gain insight into the physiological involvement of hypocretin neurons in stress-altered sleep in *5-HTT<sup>-/-</sup>* mutants versus *5-HTT<sup>+/+</sup>* mice, we performed two sets of experiments.

#### Experiment 1

RS consequences on sleep–wakefulness patterns were analyzed in *5-HTT<sup>-/-</sup>* and *5-HTT<sup>+/+</sup>* mice. To this aim, animals were restrained (RS groups) or not (control groups) for 90 min and then monitored for sleep and wakefulness. In addition, to characterize the role of hypocretin neurons in stress-altered sleep, we evaluated the consequences of hcrt-R1 blockade before RS on sleep.

#### Sleep and wakefulness analysis

**Surgery.** At 2–4 months of age, under deep sodium pentobarbital anesthesia (80 mg/kg, i.p.), animals were implanted with electrodes (made of enameled nichrome wire; diameter, 150  $\mu\text{m}$ ) for polygraphic sleep monitoring (Boutrel et al., 2002). Briefly, two electroencephalogram (EEG) electrodes were positioned onto the dura through holes made into the skull (2 mm lateral and 2 mm caudal to the bregma suture; 1 mm caudal to the lambda at midline), two electrooculogram electrodes were placed subcutaneously on each side of the orbit, and two electromyogram electrodes were inserted into the neck muscles. All electrodes were anchored to the skull with Superbond (GACD) and acrylic cement and were soldered to a miniconnector also embedded in cement. The animals were transferred to individual recording cages (20  $\times$  20  $\times$  30 cm) and were



**Figure 2.** Restraint stress alters REMS pattern in 5-HTT<sup>+/+</sup> and 5-HTT<sup>-/-</sup> mice. Mice were, or were not, restrained for 90 min (5:30 P.M. to 7:00 P.M.) and then monitored for sleep and wakefulness starting at 7:00 P.M. **A**, REMS amounts, absolute number, and mean duration of REMS episodes for the first 2 h (7:00 P.M. to 9:00 P.M. period) after the 90 min control (C) or RS session in 5-HTT<sup>+/+</sup> and 5-HTT<sup>-/-</sup> mice. **B**, REMS characteristics for the following 8 h period (11:00 P.M. to 7:00 A.M.) in the same control (C) and RS 5-HTT<sup>+/+</sup> and 5-HTT<sup>-/-</sup> mice. Data are the mean  $\pm$  SEM of independent determinations in eight animals in each group. \* $p < 0.05$ , \*\* $p < 0.01$ , significantly different from the control condition; paired Student's *t* test after ANOVA.

allowed to recover for 10 d under standard conditions (see above). They were habituated to the recording cables for 2–3 d before recordings were started.

**Recordings.** The mice were connected to the recording system with a light-weight cable and a swivel allowing free movements in the cage. Animals were recorded by groups of four, usually two 5-HTT<sup>-/-</sup> and two 5-HTT<sup>+/+</sup> mice. Spontaneous sleep–wakefulness patterns were first examined by recording mice for 24 h (baseline recordings).

**Restraint stress.** RS was performed by wrapping each mouse in a nylon-covered metal mesh during 90 min (5:30 P.M. to 7:00 P.M.), as described previously (Boutrel et al., 2002; Léna et al., 2004). At the end of RS session, mice were returned to their cage for sleep–wakefulness monitoring until 5:00 P.M. the next day. Matched control recordings were obtained the day before RS procedure, after disconnecting (5:30 P.M.) and connecting again (7:00 P.M.) the recording cables. Sleep–wakefulness was monitored for 22 h thereafter, as after the RS session.

**Pharmacological treatment.** To determine the consequences of hcrt-R1 blockade on sleep–wakefulness patterns in 5-HTT<sup>-/-</sup> and 5-HTT<sup>+/+</sup> mice, we administered the hcrt-R1 antagonist SB-334867 [1-(2-methylbenzoxazol-6-yl)-3-[1,5]naphthyridin-4-yl urea HCl] (Tocris Bioscience) at 5:00 P.M. [30 mg/kg, i.p., dissolved in 10% hydroxy-propyl- $\beta$ -cyclodextrin (Kleptose; Roquette Frères) and 1% dimethylsulfoxide]. The dose of 30 mg/kg SB-334867 was chosen because a higher dose (60 mg/kg), tested in a pilot study ( $n = 4$ ), produced in half of the animals a nonspecific state close to sedation with immobility and slightly desynchronized EEG (data not shown). In addition, at the dose of 30 mg/kg, SB-334867 was already shown to prevent hcrt-1-induced inhibition of REMS in rats (Smith et al., 2003). For baseline data, mice were injected intraperitoneally with the vehicle only. Other groups of mice received the antagonist or vehicle, before the RS or control procedure, and polygraphic recordings were collected as above.

**Scoring and analysis.** Polygraphic recordings were visually scored every 5 s epoch as wakefulness (W), non-REM sleep (NREMS), or REMS following classical criteria (Boutrel et al., 2002; Léna et al., 2004) using the Somnologia software (Medcare). NREMS latency was defined as time elapsed between the end of RS or control procedures and the first NREMS bout (lasting at least 15 s). To assess the RS effect on REMS latency, we also determined the time interval between the beginning of the first NREMS bout and the occurrence of the first REMS bout (lasting at least 15 s), after RS and control procedures. Total amount and the number and mean duration of episodes of each vigilance state were averaged over 2 h periods and over the 11:00 P.M. to 7:00 A.M. segment (8 h). Analyses were conducted using Matlab (MathWorks) routine (Léna et al., 2004).

## Experiment 2

We determined the influence of RS on serotonergic and hypocretinergic neurotransmissions in mice of both genotypes. To this aim, 5-HTT<sup>-/-</sup> and 5-HTT<sup>+/+</sup> mice were restrained (RS groups) or not (control groups) for 90 min. Serotonergic neurotransmission was evaluated by measuring 5-HT and 5-hydroxyindole acetic acid (5-HIAA) tissue contents by HPLC. Hypocretinergic neurotransmission was evaluated by determination of (1) the number of active hypocretin neurons, characterized by double preprohypocretin (preprohcrt) and c-Fos immunolabeling, (2) hcrt-1 tissue contents by radioimmunoassay, and (3) hcrt-R1 and hcrt-R2 mRNA levels by real-time PCR.

### Restraint stress

Mice were subjected to the same RS procedure as the one described above. At the end of the RS session, mice were immediately killed by decapitation or by deep anesthesia for perfusion. To evaluate the influence of RS on hypocretin neuron activation, another group of mice was returned to their home cage to be perfused 5 h later.

### Measurements of 5-HT and 5-HIAA

Restrained and control mice of each genotype were decapitated, and their brains were rapidly removed in the cold (0°C). The hypothalamus was dissected, weighed, and homogenized in 10 volumes (v/w) of 0.1 N HClO<sub>4</sub> containing 0.05% Na<sub>2</sub>S<sub>2</sub>O<sub>5</sub> and 0.05% disodium EDTA. Homogenates were centrifuged at 30,000  $\times$  g for 15 min at 4°C, and the supernatants were neutralized with 2 M KH<sub>2</sub>PO<sub>4</sub>/K<sub>2</sub>HPO<sub>4</sub>, pH 7.4, supplemented with ascorbate oxidase (final concentration, 10  $\mu$ g/ml; Boehringer Mannheim). After a second centrifugation as above to eliminate KClO<sub>4</sub> precipitate, the clear supernatants were saved, and aliquots (10–20  $\mu$ l) were injected directly into a HPLC column (25 cm, 0.46 cm outer diameter, 5  $\mu$ m particle size; Ultrasphere IP; Beckman Coulter) protected with a Brownlee precolumn (3 cm, 5  $\mu$ m). The mobile phase (at a flow rate of 1 ml/min) consisted of 70 mM KH<sub>2</sub>PO<sub>4</sub>, 2 mM triethylamine, 0.1 mM disodium EDTA, 1.25 mM octane sulfonic acid, and 16% methanol, adjusted to pH 3.02 with solid citric acid. Electrochemical detection (ESA 5011) with dual coulometric monitoring electrodes (+50 and +350 mV) was coupled with a computing integrator (System-Gold; Beckman Coulter) for the quantitative determinations of 5-HT and 5-HIAA with reference to pure standards (Hamon et al., 1988).

### Functional neuroanatomy

**Preparation of brain sections.** Animals of each phenotype were deeply anesthetized with sodium pentobarbital (80 mg/kg, i.p.) and perfused

**Table 1. Effects of pharmacological blockade of hcrt-R1 by SB-334867 (30 mg/kg, i.p.) on vigilance states during the first hour after injection in 5-HTT<sup>+/+</sup> and 5-HTT<sup>-/-</sup> mice**

	Wake		NREMS		REMS	
	5-HTT <sup>+/+</sup>	5-HTT <sup>-/-</sup>	5-HTT <sup>+/+</sup>	5-HTT <sup>-/-</sup>	5-HTT <sup>+/+</sup>	5-HTT <sup>-/-</sup>
Vehicle						
Amounts (min)	34.0 ± 2.5	32.5 ± 2.3	21.9 ± 2.2	22.5 ± 1.9	4.1 ± 0.9	5.0 ± 0.8
Episodes (absolute number)	40.6 ± 3.9	29.9 ± 2.4 <sup>#</sup>	39.9 ± 3.9	29.4 ± 2.3 <sup>#</sup>	10.0 ± 2.4	6.9 ± 1.7
SB-334867						
Amounts (min)	32.8 ± 1.8	36.0 ± 1.1	23.5 ± 1.9	21.2 ± 1.2	3.7 ± 0.7	2.7 ± 0.6
Episodes (absolute number)	41.3 ± 1.7	29.0 ± 1.4 <sup>###</sup>	40.9 ± 1.6	28.7 ± 1.3 <sup>###</sup>	9.6 ± 2.7	6.4 ± 1.1

Values are expressed as the mean ± SEM of seven animals in each group. <sup>#</sup>*p* < 0.05, <sup>###</sup>*p* < 0.001, 5-HTT<sup>-/-</sup> mutants versus wild-type mice; Fisher's PLSD test after ANOVA.

**Table 2. Effects of acute blockade of hcrt-R1 by SB-334867 on vigilance state amounts during the first 2 h after restraint stress in 5-HTT<sup>+/+</sup> and 5-HTT<sup>-/-</sup> mice**

	Wake (min)		NREMS (min)		REMS (min)	
	5-HTT <sup>+/+</sup>	5-HTT <sup>-/-</sup>	5-HTT <sup>+/+</sup>	5-HTT <sup>-/-</sup>	5-HTT <sup>+/+</sup>	5-HTT <sup>-/-</sup>
Vehicle + control (day 1)	76.3 ± 3.5	73.9 ± 5.4	38.3 ± 3.3	39.6 ± 5.1	5.4 ± 0.7	6.5 ± 0.6
Vehicle + stress (day 2)	74.8 ± 4.7	68.0 ± 4.8	42.3 ± 4.8	48.8 ± 4.3	2.9 ± 0.4 <sup>**</sup>	3.2 ± 1.0 <sup>***</sup>
Vehicle + control (day 1)	68.9 ± 3.3	64.9 ± 5.4	43.3 ± 2.7	45.6 ± 4.7	7.8 ± 1.0	9.5 ± 1.5
SB-334867 + stress (day 2)	64.2 ± 5.8	68.6 ± 2.8	51.6 ± 6.2	47.6 ± 2.8	4.2 ± 1.1 <sup>*</sup>	3.9 ± 0.6 <sup>*</sup>

See Figure 3A for the experimental protocol. Values are expressed as the mean ± SEM of eight to nine mice in each group. <sup>\*</sup>*p* < 0.05, <sup>\*\*</sup>*p* < 0.01, and <sup>\*\*\*</sup>*p* < 0.001, restraint stress versus control condition; paired Student's *t* test after ANOVA.

intracardially with 100 ml of 4% paraformaldehyde (PFA) in PBS (50 mM NaH<sub>2</sub>PO<sub>4</sub>/Na<sub>2</sub>HPO<sub>4</sub> and 154 mM NaCl, pH 7.4). Brains were removed, immersed for 12 h in the same fixative, and cryoprotected for 24 h in 30% sucrose in PBS. Brains were then quickly frozen in isopentane chilled at -30°C with dry ice and stored at -20°C. The frozen brains were cut into three series of 30-μm-thick coronal sections at the level of the hypothalamic perifornical region using a cryostat (CM3050S; Leica). The collected sections were stored at -20°C in cryoprotectant solution (30% glycerol and 30% ethylene glycol in PBS).

#### Double immunohistochemistry

**Hypocretin/c-Fos.** One series of sections was used to detect both c-Fos and preprohcrt in the hypothalamic perifornical region. For the first step (c-Fos immunostaining), sections were incubated overnight at 4°C with rabbit polyclonal anti-c-Fos antiserum [1:20,000, c-Fos (ab-5); Oncogene Science] in PBS containing 4% bovine serum albumin (BSA) and 0.1% Triton X-100. After rinsing in PBS supplemented with 0.1% Triton X-100, sections were incubated for 2 h with biotinylated anti-rabbit IgG (1:200 dilution with ABC Vectastain Elite kit; Vector Laboratories) in PBS containing 4% BSA and 0.1% Triton X-100. Additional rinses were followed by incubation with avidin–biotin–horseradish peroxidase solution (ABC Vectastain Elite kit; Vector Laboratories) for 1 h, and c-Fos immunoreactivity was visualized as a black reaction product after incubation of the sections in 0.05 M Tris-HCl, pH 7.6, containing 0.04% (w/v) of 3,3'-diaminobenzidine (DAB) (Sigma) and 0.2% ammonium nickel sulfate (Sigma), supplemented with increasing concentrations of H<sub>2</sub>O<sub>2</sub> (from 0.00015 to 0.0048%). For the second step, sections were rinsed and incubated overnight with a rabbit anti-preprohcrt antiserum (1:200 dilution; Alpha Diagnostics). After rinses, sections were treated with 0.03% H<sub>2</sub>O<sub>2</sub> in PBS to inactivate remaining peroxidase activity. The immunohistochemical procedure was then processed as described previously. For staining, sections were treated with 0.04% DAB in PBS, yielding a brown cytoplasmic product in preprohcrt-immunoreactive (IR) neurons. Stained sections were finally mounted on gelatin-coated slides, dehydrated, and coverslipped with Eukitt mounting medium (Kindler).

**Cell counting.** The resultant signals were quantified using the cell counting software Mercator Lite 3.0B (Explora Nova). Two examiners, blind to the experimental conditions, performed all the cell counts using a microscope at various magnifications (2.5× to 40×), as needed. Double-labeled neurons were carefully analyzed at high magnification (40×). All perifornical preprohcrt-IR and double-labeled neurons were counted on both sides of the brain in two different sections, 90 μm apart, without considering the intensity of the staining. For each mouse, the number of labeled cells of each type was then averaged per section.

#### Hcrt-1 tissue contents

**Peptide extraction.** Restrained and control mice of each genotype were decapitated, and the anterior part of the raphe area (which contains 5-HT cell bodies of the median and dorsal raphe nuclei) was dissected out and frozen at -80°C until use. For peptide extraction, frozen tissues were homogenized by sonication in 100 μl of 0.1 N HCl and boiled for 15 min. After centrifugation (30,000 × *g*, 15 min, 4°C), the supernatant was adjusted to pH 7.0 with 1 M Tris base. The resulting precipitate was spun down by a second centrifugation at 30,000 × *g* for 15 min at 4°C. The clear extract was assayed for its hcrt-1 content at 1:10 dilution, and residues were allotted to protein assays.

**Hcrt-1 assay.** Hcrt-1 contents were measured using a [<sup>125</sup>I]RIA kit (Phoenix Pharmaceuticals). Briefly, peptide extracts were diluted in RIA buffer, and duplicate samples were assayed. Hcrt-1 levels were determined against a known standard established with known quantities of pure hcrt-1.

**Protein assay.** Proteins were quantified using the Folin's method (Lowry et al., 1951) with BSA as standard.

#### hcrt-R1 and hcrt-R2 mRNA expression

**RNA extraction and first-stranded cDNA synthesis.** Mice from each genotype were killed by decapitation, and the entire brain was removed. The anterior raphe area was dissected, frozen in liquid nitrogen, and stored at -80°C until use. Total RNA extraction was done using the Stratagene kit (Absolutely RNA Purification kit), according to the instructions of the manufacturer, including removal of genomic DNA by DNase treatment (Stratagene). RNA integrity was checked by agarose gel electrophoresis, and RNA concentrations were determined by spectrophotometric measurements. First-stranded cDNA synthesis (1 μg of total RNA per 20 μl reaction) was performed with the SuperScript III Reverse Transcriptase and random primers, as recommended by the manufacturer (Invitrogen).

**Quantitative TaqMan reverse transcription-PCR.** PCR amplification in triplicate for each sample was performed (ABI Prism 7300; Applied Biosystems) by using ABsolute QPCR Mixes (ABgene) and the Assays-on-Demand Gene Expression probes (Applied Biosystems) for hcrt-R1 and hcrt-R2 mRNAs and mRNAs from reporter genes encoding glyceraldehyde 3-phosphate dehydrogenase (GAPDH) and hypoxanthine guanine phosphoribosyltransferase (HPRT). The polymerase activation step at 95°C for 15 min was followed by 40 cycles of 15 s at 95°C and 60 s at 60°C. The validity of the results was checked by running appropriate negative controls (substitution of cDNA with water for PCR amplification or omission of reverse transcriptase for the cDNA synthesis step).

**Data analysis.** The 2<sup>-ΔΔCt</sup> method (Livak and Schmittgen, 2001) was used to analyze the relative change in gene expression in 5-HTT<sup>-/-</sup> and



$5\text{-HTT}^{+/+}$  mice under control condition or after RS (RQ Study Software version 1.2; Applied Biosystems). All data (expressed in arbitrary units) were analyzed using both GAPDH and HPRT as references.

### Statistical analyses

For experiment 1, RS consequences on the sleep–wakefulness patterns (amount, absolute number, and mean duration of each vigilance state) were first analyzed by a two-way (genotype and stress) ANOVA, with repeated measures over time (states of vigilance over 2 h segments). When appropriate, additional two-way ANOVAs were performed for stress (repeated measures, RS vs control) and genotype ( $5\text{-HTT}^{-/-}$  vs  $5\text{-HTT}^{+/+}$ ) on specific time slots. The influence of hcrt-R1 pharmacological blockade was assessed by a two-way ANOVA for treatment (repeated measures, SB-334867 vs vehicle) and genotype ( $5\text{-HTT}^{-/-}$  vs  $5\text{-HTT}^{+/+}$ ). For pharmacological treatments combined with RS, data were analyzed independently for mutant and wild-type mice using analyses of covariance (ANCOVAs) to assess the influence of SB-334867 on stress response. The ANCOVAs were performed on W, NREMS, or REMS amounts after RS as dependent variable, pharmacological treatment (SB-334867 vs vehicle) as fixed factor, and the amount of vigilance state before stress as covariate. When appropriate, ANOVAs and ANCOVAs were followed by pairwise comparisons using unpaired or paired  $t$  tests as needed. The level of significance was set at  $p < 0.05$ .

For experiment 2, data were analyzed using a two-way (genotype and stress) ANOVA, followed as appropriate by Fisher's protected least significant difference (PLSD) test for pairwise multiple comparisons.

## Results

No significant differences were found between mice obtained from homozygous or heterozygous breeding under basal conditions, as well as after RS. Thus, the data were pooled independently of the breeding condition.

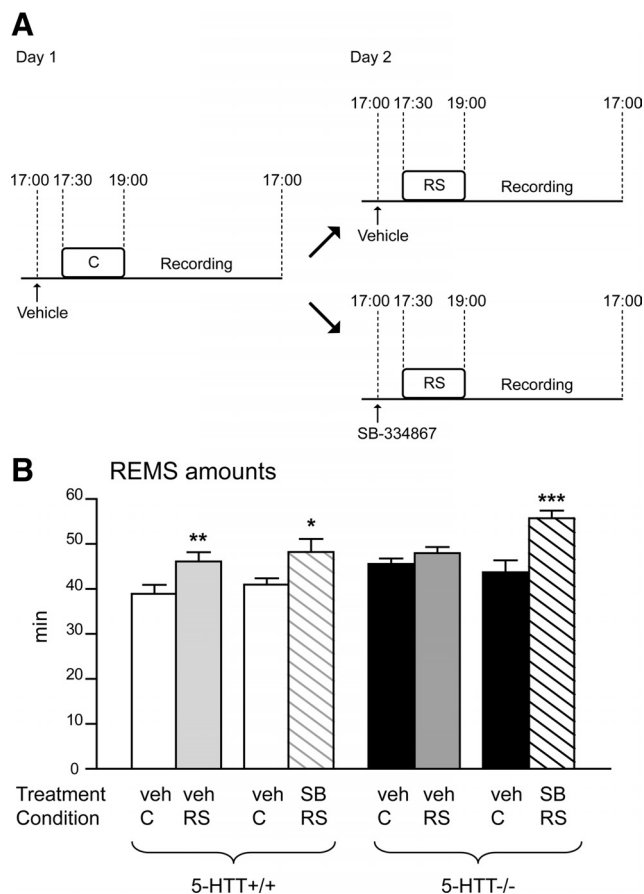
### Stress-altered sleep in $5\text{-HTT}^{-/-}$ and $5\text{-HTT}^{+/+}$ mice: influence of the hypocretinergic system

#### Sleep–wakefulness patterns after restraint stress in $5\text{-HTT}^{-/-}$ and $5\text{-HTT}^{+/+}$ mice

As described previously (Wisor et al., 2003; Alexandre et al., 2006),  $5\text{-HTT}^{-/-}$  mice exhibited, at baseline, a 45% increase in REMS amounts compared with wild-type mice ( $F_{(1,18)} = 17.89$ ,  $p < 0.001$ , over 24 h; data not shown), which was accounted for by a 43% increase in the number of REMS episodes ( $F_{(1,18)} = 12.20$ ,  $p < 0.01$ ). This increase was mainly observed during the light period (data not shown). However, REMS enhancement in mutant mice was attenuated in our control conditions [after disconnecting (5:30 P.M.) and connecting again (7:00 P.M.) the recording cables], because it reached only +19% over REMS amounts in wild-type mice ( $F_{(1,14)} = 6.40$ ,  $p < 0.05$ ), suggesting an increased (stress) reactivity to manipulation in  $5\text{-HTT}^{-/-}$  mice.

The most remarkable changes elicited by 90 min RS on sleep–wakefulness patterns concerned REMS characteristics. Although RS induced short-term changes in REMS that were similar in both genotypes, this stress session affected differently REMS in  $5\text{-HTT}^{-/-}$  and  $5\text{-HTT}^{+/+}$  mice during late recovery (Fig. 1).

First, RS significantly lengthened REMS latency in both genotypes (stress,  $F_{(1,14)} = 22.35$ ,  $p < 0.001$ ; in minutes, control conditions,  $22.4 \pm 12.7$  and  $12.0 \pm 2.3$  vs RS,  $77.9 \pm 13.1$  and  $75.1 \pm 24.2$ , in  $5\text{-HTT}^{+/+}$  and  $5\text{-HTT}^{-/-}$  mice, respectively), although it did not modify the NREMS latency. Thus, mice of both genotypes exhibited a significant reduction of REMS amounts during the first 2 h after RS ( $p < 0.05$ ) (Figs. 1, 2A). This change was accounted for by a decrease in the number of REMS episodes (stress,  $F_{(1,14)} = 13.23$ ,  $p < 0.01$ ), with no modification in their mean duration (Fig. 2A). After this 2 h period,



**Figure 3.** Effects of acute treatment with the hcrt-R1 antagonist SB-334867 before restraint stress on sleep–wakefulness patterns in  $5\text{-HTT}^{+/+}$  and  $5\text{-HTT}^{-/-}$  mice. **A**, Experimental protocol: on the first day of recording (Day 1),  $5\text{-HTT}^{+/+}$  and  $5\text{-HTT}^{-/-}$  mice were injected with the vehicle at 5:00 P.M. and then subjected to the control (C) procedure for stress by disconnecting (5:30 P.M.) and connecting again (7:00 P.M.) the recording cable to the mouse. Recordings were obtained until 5:00 P.M. the next day. On the following day (Day 2), at 5:00 P.M.,  $5\text{-HTT}^{+/+}$  and  $5\text{-HTT}^{-/-}$  mice were injected with the vehicle or the hcrt-R1 antagonist SB-334867 (30 mg/kg, i.p.), and RS was started 30 min later (at 5:30 P.M.). Recordings of vigilance states were performed as for the preceding control step (Day 1). **B**, REMS amounts for 8 h starting at 11:00 P.M. under control conditions (C) or after RS in  $5\text{-HTT}^{+/+}$  and  $5\text{-HTT}^{-/-}$  mice previously injected with the vehicle (veh) or SB-334867 (SB). Data are the mean  $\pm$  SEM of eight to nine animals in each group. \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ , significantly different from the respective control condition; paired Student's  $t$  test after ANOVA.

**Table 3. Effects of restraint stress on 5-HT and 5-HIAA tissue levels in the hypothalamus of  $5\text{-HTT}^{+/+}$  and  $5\text{-HTT}^{-/-}$  mice**

	$5\text{-HTT}^{+/+}$		$5\text{-HTT}^{-/-}$	
	Control	RS	Control	RS
$n$	7	8	7	8
5-HT (ng/g tissue)	$959 \pm 30$	$971 \pm 49$	$205 \pm 23^{***}$	$265 \pm 26^{***}$
5-HIAA (ng/g tissue)	$394 \pm 13$	$452 \pm 26$	$220 \pm 17^{***}$	$333 \pm 31^{***,*}$
5-HIAA/5-HT	$0.41 \pm 0.01$	$0.47 \pm 0.03$	$1.08 \pm 0.08^{***}$	$1.26 \pm 0.05^{***,*}$

Values are expressed as the mean  $\pm$  SEM of the number of animals indicated ( $n$ ). \*\* $p < 0.01$ , \*\*\* $p < 0.001$ ,  $5\text{-HTT}^{-/-}$  mutants versus wild-type mice; Fisher's PLSD test after ANOVA.

\* $p < 0.05$  and \*\* $p < 0.01$ , restraint stress (RS) versus control condition; Fisher's PLSD test after ANOVA.

RS affected differently REMS amounts across genotypes (genotype  $\times$  stress interaction,  $F_{(1,294)} = 8.63$ ,  $p < 0.01$ ) (Fig. 1). Indeed, the initial drop was followed by an increase in REMS amounts in  $5\text{-HTT}^{+/+}$  mice (stress,  $F_{(1,14)} = 20.18$ ,  $p < 0.001$ ; during the 11:00 P.M. to 7:00 A.M. period) (Fig. 2B) but not in  $5\text{-HTT}^{-/-}$  mutants (genotype  $\times$  stress interaction,  $F_{(1,14)} = 8.74$ ,

**Table 4. Restraint stress-induced activation of hypocretineric neurons in *5-HTT*<sup>+/+</sup> and *5-HTT*<sup>-/-</sup> mice**

	<i>t</i> <sub>0</sub>				<i>t</i> <sub>0</sub> + 5 h			
	<i>5-HTT</i> <sup>+/+</sup>		<i>5-HTT</i> <sup>-/-</sup>		<i>5-HTT</i> <sup>+/+</sup>		<i>5-HTT</i> <sup>-/-</sup>	
	Control	RS	Control	RS	Control	RS	Control	RS
<i>n</i>	7	9	8	8	8	7	8	8
Preprohcrt-IR positive neurons	307 ± 21	305 ± 17	296 ± 11	292 ± 22	265 ± 12	287 ± 16	285 ± 11	267 ± 5
Preprohcrt-IR/c-Fos-IR positive neurons	61 ± 19	170 ± 11***	56 ± 18	156 ± 14***	153 ± 9	203 ± 10***	167 ± 12	216 ± 5***

The data are given as number of labeled neurons per section in the perifornical region of wild-type and mutant mice. For each mouse, among all the observed sections of the perifornical region, the two most intensively labeled sections were analyzed. Control and stressed mice were killed immediately (*t*<sub>0</sub>) or 5 h (*t*<sub>0</sub> + 5 h) after RS or control conditions. Values are expressed as the mean ± SEM of the number of animals indicated (*n*). \*\*\**p* < 0.001, RS versus control condition; Fisher's PLSD test after ANOVA.

*p* < 0.05) (Figs. 1, 2*B*). This REMS increase in wild-type mice was accounted for by an enhancement in the number of episodes, with no change in their mean duration (Fig. 2*B*).

In addition, RS induced changes in W and NREMS amounts in both strains (stress; W,  $F_{(1,294)} = 12.70$ , *p* < 0.001; NREMS,  $F_{(1,294)} = 15.16$ , *p* < 0.001). More specifically, stressed *5-HTT*<sup>+/+</sup> mice showed a decrease in W (-18%; *p* < 0.001) and an increase in NREMS (+21%; *p* < 0.001) amounts during the 12 h of the dark period (Fig. 1 and data not shown). Similar trends were observed in *5-HTT*<sup>-/-</sup> mice, but corresponding changes were of lower amplitude (W, -9%; NREMS, +10%; not significant) (Fig. 1).

Altogether, these data show that *5-HTT*<sup>-/-</sup> mice exhibited attenuated responses to restraint stress and notably an absence of secondary REMS enhancement. In contrast, on a short-term basis, restraint stress inhibited REMS (reduced amount and number of episodes; increased latency) independently of the mouse genotype.

#### Acute hcrt-R1 blockade in *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice

Because hypocretins exert a strong inhibitory influence on REMS, we investigated whether hypocretineric neurotransmission contributed, at least in part, to the effects of RS on sleep-wakefulness patterns in *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice. To test this hypothesis, we blocked hcrt-R1 in mice at the time of RS.

**Effects of SB-334867 under basal conditions.** We first evaluated the consequences of acute blockade of hcrt-R1 with the selective antagonist SB-334867 on sleep and wakefulness patterns in non-stressed *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice. As observed previously in rats (Smith et al., 2003) and mice (Adamantidis et al., 2007), SB-334867 (30 mg/kg, i.p.) did not significantly affect W, NREMS, or REMS amounts during the first hour (Table 1) and the 12 h (data not shown) after injection in both wild-type and *5-HTT*<sup>-/-</sup> mice. In addition, transitions from sleep to wake were also unchanged by SB-334867 in both strains (data not shown).

**Effects of SB-334867 under restraint stress conditions (see protocol in Fig. 3*A*).** The influence of stress on W and NREMS amounts was not significantly affected by previous administration of SB-334867 in both *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice [first 2 h after restraint stress (Table 2); 11:00 P.M. to 7:00 A.M.; data not shown].

However, pharmacological blockade of hcrt-R1 had contrasting influences on REMS in the two strains, regarding notably the delayed influences of RS. Thus, injection of SB-334867 30 min before RS (at 5:00 P.M.) (Fig. 3*A*) induced in *5-HTT*<sup>-/-</sup> mice a significant increase in REMS amounts during the 11:00 P.M. to 7:00 A.M. period (ANCOVA,  $F_{(1,13)} = 14.95$ , *p* < 0.01) (Fig. 3*B*). Conversely, in wild-type mice, REMS enhancement after RS was occurring similarly with or without previous pharmacological blockade of hcrt-R1 by SB-334867 (ANCOVA for treatment,

$F_{(1,14)} = 0.14$ , *p* = 0.72) (Fig. 3*B*). It should be noted, however, that *5-HTT*<sup>+/+</sup> mice exhibited an attenuated stress-altered sleep after treatment with vehicle compared with naive animals, notably for the REMS increase (Fig. 3*B* vs Fig. 2*B*). This could have been a consequence of the experimental paradigm that imposed vehicle injection on the control day (control condition; see Materials and Methods). As observed previously, the dramatic decrease in REMS amounts during the first 2 h after RS occurred in both genotypes independently of the pharmacological treatment (vehicle group,  $F_{(1,15)} = 47.87$ , *p* < 0.0001; SB-334867 group,  $F_{(1,14)} = 16.04$ , *p* < 0.01) (Table 2).

Therefore, in *5-HTT*<sup>-/-</sup> mice, the pharmacological blockade of hcrt-R1 before stress restored the delayed increase in REMS amounts that was regularly observed in *5-HTT*<sup>+/+</sup> mice after RS.

#### Effects of restraint stress on serotonergic neurotransmission in *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice

##### *5-HT* and *5-HIAA* tissue levels after restraint stress in *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice

Stress has been reported to excite serotonergic neurons in rats and mice (Chaouloff, 1993; Cullinan et al., 1995; Grahn et al., 1999; Roche et al., 2003). Accordingly, we examined whether 90 min RS affected differentially 5-HT and 5-HIAA tissue levels in *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice by using HPLC.

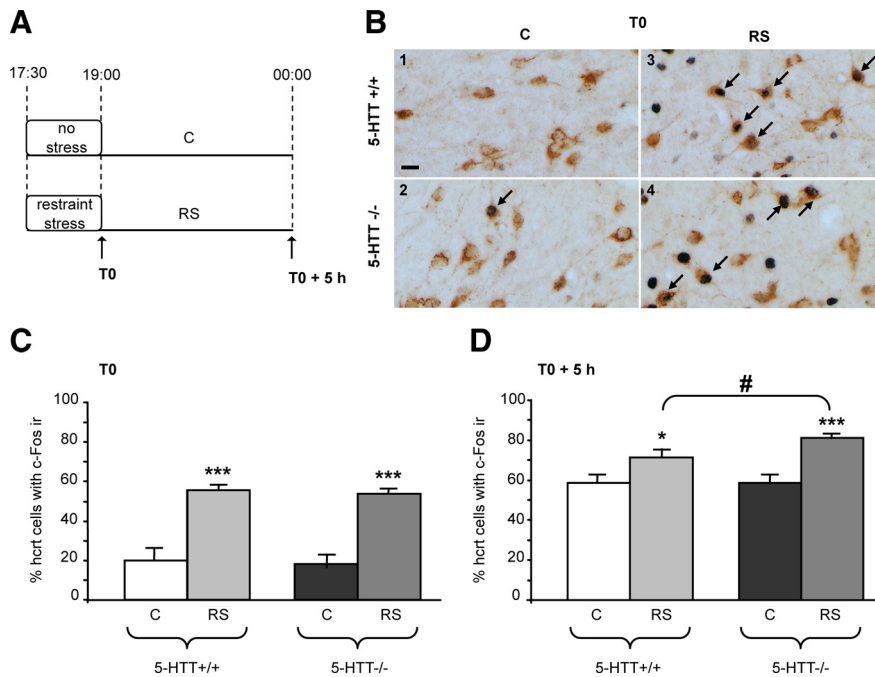
As described previously in several forebrain structures (Fabre et al., 2000; Kim et al., 2005), 5-HT levels in the hypothalamus of *5-HTT*<sup>-/-</sup> mutants were strongly decreased (approximately -80% compared with *5-HTT*<sup>+/+</sup> mice) both under control condition and after RS ( $F_{(1,26)} = 449.04$ , *p* < 0.001) (Table 3). A less pronounced decrease (-26 to -40%) was observed for 5-HIAA levels ( $F_{(1,26)} = 37.44$ , *p* < 0.001) (Table 3). These genotype-dependent changes resulted in marked increases of 5-HIAA/5-HT ratios in *5-HTT*<sup>-/-</sup> mutants (approximately +160% over those in *5-HTT*<sup>+/+</sup> mice;  $F_{(1,26)} = 236.41$ , *p* < 0.001) (Table 3).

In *5-HTT*<sup>-/-</sup> mice, RS was found to increase significantly the tissue levels of 5-HIAA in the hypothalamus (+51%;  $F_{(1,26)} = 12.76$ , *p* < 0.01) (Table 3) and nonsignificantly those of 5-HT ( $F_{(1,26)} = 1.12$ , *p* = 0.30). RS also increased significantly the 5-HIAA/5-HT ratio (+17% in the mutants). In contrast, these changes did not reach statistical significance in *5-HTT*<sup>+/+</sup> mice (Table 3). Accordingly, RS activated the hypothalamic serotonergic neurotransmission to a higher degree in *5-HTT*<sup>-/-</sup> mutants than in *5-HTT*<sup>+/+</sup> mice.

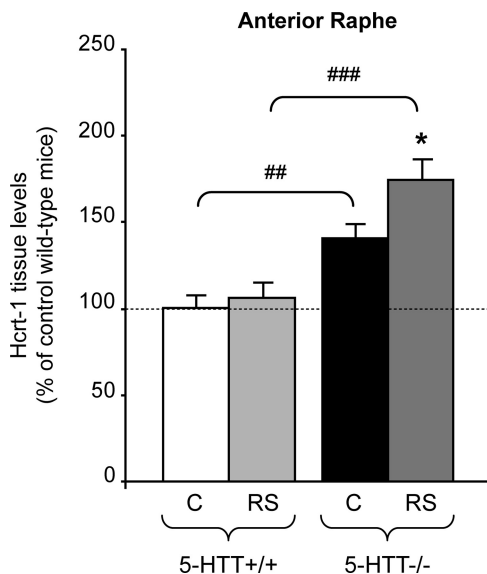
#### Effects of restraint stress on hypocretineric neurotransmission in *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice

##### Activation of hypocretin neurons

Stress has also been reported to excite hypocretin neurons in rats and mice (Estabrooke et al., 2001; Espana et al., 2003; Winsky-Sommerer et al., 2004). We therefore examined the effects of a 90



**Figure 4.** Activation of hypocretin neurons after restraint stress in *5-HTT*<sup>+/+</sup> and *5-HTT*<sup>-/-</sup> mice. **A**, Schematic representation of the experimental protocol. The activation of hypocretin neurons has been measured in *5-HTT*<sup>+/+</sup> and *5-HTT*<sup>-/-</sup> mice immediately ( $t_0$ ) or 5 h ( $t_0 + 5$  h) after the end of 90 min RS and in nonstressed (control; C) paired mice killed at the same time. **B**, Representative immunocytochemical labeling at the level of the perifornical area of the lateral hypothalamus showing few hypocretin-IR cells (brown staining) with c-Fos-IR labeling (dark staining) in control conditions in *5-HTT*<sup>+/+</sup> (1) and *5-HTT*<sup>-/-</sup> (2) mice. RS induced a strong and immediate ( $t_0$ ) increase in the number of c-Fos-IR nuclei in hypocretin-IR neurons in both *5-HTT*<sup>+/+</sup> (3) and *5-HTT*<sup>-/-</sup> (4) mice. Black arrows, c-Fos-IR and hypocretin-IR neurons. Scale bar, 10  $\mu$ m. **C, D**, Percentage of hypocretin-IR neurons with c-Fos-IR labeling in control conditions (C) and after RS in *5-HTT*<sup>+/+</sup> and *5-HTT*<sup>-/-</sup> mice. Mice were killed immediately ( $t_0$ ; **C**) or 5 h ( $t_0 + 5$  h; **D**) after RS or control conditions. Data are expressed as the mean  $\pm$  SEM of seven to eight mice in each group. \* $p < 0.05$ , \*\*\* $p < 0.001$ , significantly different from the control condition; # $p < 0.05$ , *5-HTT*<sup>-/-</sup> versus wild-type mice; Fisher's PLSD test after ANOVA.



**Figure 5.** Effects of restraint stress on hcr1 levels in the anterior raphe area of *5-HTT*<sup>+/+</sup> and *5-HTT*<sup>-/-</sup> mice. Hcr1 tissue levels were measured in the anterior raphe area of *5-HTT*<sup>+/+</sup> and *5-HTT*<sup>-/-</sup> mice in control condition (C) or immediately after RS. Hcr1 tissue levels are expressed as percentage of values in nonstressed *5-HTT*<sup>+/+</sup> mice. Absolute values were as follows (ng/ $\mu$ g protein): 1.46  $\pm$  0.11 (control condition in *5-HTT*<sup>+/+</sup>), 1.54  $\pm$  0.13 (RS in *5-HTT*<sup>+/+</sup>), 2.09  $\pm$  0.11 (control condition in *5-HTT*<sup>-/-</sup>), and 2.56  $\pm$  0.16 (RS in *5-HTT*<sup>-/-</sup>). Data are the mean  $\pm$  SEM of seven to nine animals in each group. \* $p < 0.05$ , significantly different from the control condition; # $p < 0.01$ , ### $p < 0.001$ , *5-HTT*<sup>-/-</sup> versus wild-type mice; Fisher's PLSD test after ANOVA.

min RS session on hypocretin neuron activity in *5-HTT*<sup>-/-</sup> versus *5-HTT*<sup>+/+</sup> mice by using the c-Fos marker.

Under control conditions (at 7:00 P.M.; control  $t_0$ ), *5-HTT*<sup>-/-</sup> mice showed no differences in the total number of hypocretin neurons in the perifornical region of the hypothalamus compared with wild-type mice (Table 4). In both strains, the number of hypocretin neurons expressing c-Fos was markedly increased (by  $\sim 2.8$ -fold) immediately after RS (RS  $t_0$ ) ( $F_{(1,28)} = 45.77$ ,  $p < 0.001$ ) (see Fig. 5B, Table 4), whereas the total number of hypocretin neurons remained unchanged ( $F_{(1,28)} = 0.02$ ,  $p = 0.88$ ) (Table 4). As a consequence, stressed mice of both genotypes exhibited a greater percentage of double-labeled neurons than respective unstressed controls ( $F_{(1,28)} = 65.60$ ,  $p < 0.001$ ) (Fig. 4C). Furthermore, RS activated hypocretin neurons to the same degree in both strains (Fig. 4C).

To determine the time course of hypocretin neuron activation after RS, some mutant and wild-type mice were returned to their home cage after RS and were anesthetized for intracardiac perfusion with PFA at midnight ( $t_0 + 5$  h).

As shown previously in rats (Estabrooke et al., 2001), hypocretin neuron activity varied across the light/dark cycle, with higher levels of activity detected during the dark period as expected from wake-active neurons (M. G. Lee et al., 2005; Mileykovskiy et al., 2005; Takahashi et al., 2008).

Accordingly, we observed a robust increase (by approximately threefold) in the percentage of c-Fos-expressing hypocretin neurons in nonstressed control mice of both genotypes killed during the dark period (0:00, control  $t_0 + 5$  h) compared with mice killed at the end of the light period (7:00 P.M., control  $t_0$ ;  $F_{(1,27)} = 66.78$ ,  $p < 0.001$ ). However, the total number of hypocretin neurons was not significantly different at  $t_0 + 5$  h compared with  $t_0$  in both *5-HTT*<sup>+/+</sup> and *5-HTT*<sup>-/-</sup> mice (Table 4).

Five hours after restraint stress (RS  $t_0 + 5$  h), the number of hypocretin neurons expressing c-Fos was still increased in both strains compared with their respective nonstressed controls (stress,  $F_{(1,27)} = 27.51$ ,  $p < 0.001$ ) (Table 4). Accordingly, the percentage of double-labeled neurons was increased by RS in both strains (stress effect,  $F_{(1,27)} = 26.99$ ,  $p < 0.001$ ) (Fig. 4D). Interestingly, this RS effect was almost twice as large in *5-HTT*<sup>-/-</sup> mutants (+38.1%) as in *5-HTT*<sup>+/+</sup> mice (+22.5%).

In summary, RS produced an immediate and a long-lasting activation of hypocretin neurons in both *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice. However, this long-lasting influence of RS was larger in *5-HTT*<sup>-/-</sup> mice.

#### Hcr1 tissue levels after restraint stress in the anterior raphe area of *5-HTT*<sup>-/-</sup> and *5-HTT*<sup>+/+</sup> mice

Hypocretinergic neurons project to the anterior raphe nuclei (Peyron et al., 1998; H. S. Lee et al., 2005; Wang et al., 2005). Thus, hcr1 tissue levels were measured, by radioimmunoassay, in the anterior raphe area, which includes 5-HT cell bodies from



**Table 5. Effects of restraint stress on hcrt-R1 and hcrt-R2 mRNA levels in the anterior raphe area of *5-HTT*<sup>+/+</sup> and *5-HTT*<sup>-/-</sup> mice**

	<i>5-HTT</i> <sup>+/+</sup>		<i>5-HTT</i> <sup>-/-</sup>	
	Control	RS	Control	RS
<i>n</i>	7	6–7	7	7
hcrt-R1 mRNA (A.U.)	0.79 ± 0.07	1.21 ± 0.20* (+53%)	0.68 ± 0.08	1.14 ± 0.15* (+68%)
hcrt-R2 mRNA (A.U.)	1.49 ± 0.17	1.43 ± 0.17 (-4%)	1.41 ± 0.05	1.45 ± 0.11 (+3%)

Values [relative quantity, mean ± SEM of the number of animals indicated (*n*)] are expressed in arbitrary units (A.U.) after normalization to an endogenous reference gene (GAPDH). Normalization with HPRT as endogenous reference gene (see Materials and Methods) gave similar results. \**p* < 0.05, RS versus control condition; Fisher's PLSD test after ANOVA.

the dorsal and the median raphe nuclei. The two-way ANOVA indicates overall effects of genotype ( $F_{(1,26)} = 34.38$ ,  $p < 0.001$ ) and stress ( $F_{(1,26)} = 4.71$ ,  $p < 0.05$ ). As illustrated in Figure 5, under control conditions, hcrt-1 tissue levels were significantly higher in *5-HTT*<sup>-/-</sup> mutants compared with wild-type mice. Although RS did not affect hcrt-1 levels in *5-HTT*<sup>+/+</sup> mice, a significant increase was observed in stressed *5-HTT*<sup>-/-</sup> mutants (+25% over nonstressed control *5-HTT*<sup>-/-</sup> mice). As a result, after stress, hcrt-1 levels were 66% higher ( $p < 0.001$ ) in *5-HTT*<sup>-/-</sup> mutants than in paired *5-HTT*<sup>+/+</sup> mice (Fig. 5).

#### *hcrt-R1 and hcrt-R2 mRNA levels*

To test whether changes in 5-HT neurotransmission and/or RS may influence hcrt receptor expression, hcrt-R1 and hcrt-R2 mRNA levels were determined in the anterior raphe area after 90 min RS and under control conditions.

As reported in Table 5, the levels of hcrt-R1 mRNA in the anterior raphe area were not different in *5-HTT*<sup>-/-</sup> and wild-type mice under control conditions ( $F_{(1,23)} = 0.47$ ,  $p = 0.50$ ). An RS-induced increase in hcrt-R1 mRNA levels was observed in both *5-HTT*<sup>-/-</sup> (+68%) and wild-type (+53%) mice ( $F_{(1,23)} = 11.83$ ,  $p < 0.01$ ) (Table 5). In contrast, in all the conditions examined, hcrt-R2 mRNA expression remained unchanged (Table 5).

The RS-induced increase in hcrt-R1 mRNA expression in the anterior raphe area suggests that hypocretinergic neurotransmission may be enhanced in *5-HTT*<sup>-/-</sup> and wild-type mice in response to RS.

## Discussion

The present data show that genetically driven loss of the 5-HTT function produces impaired sleep modulations by stress. These impairments appeared to be causally related to an enhanced hypocretinergic neurotransmission because the sleep response to stress in *5-HTT*<sup>-/-</sup> mutant mice could be normalized by blocking the hypocretin type 1 receptor. These results emphasize the role of hypocretin neurons in the disrupted stress response consecutive to 5-HTT deficiency in mice.

### Mutant mice lacking the 5-HTT show an altered sleep homeostasis after restraint stress

It is generally agreed that sleep quality and duration depend on events taking place during wakefulness. In rodents, stress induces a marked arousal and has subsequently sleep-promoting influences that mainly concern REMS (Rampin et al., 1991; Marinesco et al., 1999; Boutrel et al., 2002). In the present study, 90 min RS session induced strong adaptations of the sleep–wakefulness patterns in wild-type male mice of the CD-1 background as described previously in CD1 females (Popa et al., 2008) and other mouse strains (Meerlo et al., 2001; Boutrel et al., 2002; Léna et al., 2004): initially, a dramatic decrease in REMS amounts that lasted for 2 h after RS, and, secondarily, an increase in REMS and NREMS amounts at the expense of wakefulness.

Compared with wild-type mice, *5-HTT*<sup>-/-</sup> mutants displayed at baseline larger REMS amounts as shown previously (Wisor et

al., 2003; Alexandre et al., 2006), but they exhibited a blunted sleep response to stress that was mainly characterized by the absence of the secondary REMS increase. This suggests that there is no linear relation between spontaneous and stress-triggered REMS amounts, in agreement with other models (Popa et al., 2008). The absence of the delayed increase in REMS after RS, which had been also observed in female *5-HTT*<sup>-/-</sup> mice (Popa et al., 2008), was associated here with a higher RS-induced activation of serotonergic neurons in *5-HTT*<sup>-/-</sup> mutants compared with wild-type mice (Table 3). Previous studies already showed that 5-HT neurons are critical for mediating the delayed influence of stress on sleep. Thus, in rats, the 5-HT outflow in the hypothalamus is strongly enhanced after RS, and mimicking such enhancement by direct DR electrical stimulation is sufficient to produce a delayed REMS increase (Houdouin et al., 1991a,b). However, because the initial decrease in REMS amounts after RS was maintained in *5-HTT*<sup>-/-</sup> mice, it can be inferred that only secondary changes in REMS are controlled by 5-HT neurotransmission. Interestingly, the inhibition of 5-HT synthesis (Sinha, 2006) or the absence of 5-HT<sub>1A</sub> or 5-HT<sub>1B</sub> receptors (Boutrel et al., 2002; Adrien et al., 2004) also abolishes the REMS induction by RS.

Our findings in *5-HTT*<sup>-/-</sup> mice are in line with previous data indicating that these mutants exhibit a number of stress-related phenotypical changes: enhanced neuroendocrine responses to stress either on the CD-1 (Li et al., 1999) or the C57BL/6J (Tjurmina et al., 2002) background and exaggerated behavioral responses to mild stressor, including saline injections (Li et al., 1999; Tjurmina et al., 2002). In addition, *5-HTT*<sup>-/-</sup> mice present an exacerbated depression-like behavior after repeated exposure to forced swim stress (Wellman et al., 2007) and enhanced sensitivity to a single predator exposure, which has been proposed as a model for post-traumatic stress disorders (Adamec et al., 2006). Thus, the impaired sleep homeostasis in response to stress found here in these mutants is consistent with the hypothesis that 5-HTT deficiency leads to alterations in the capacity to adapt to environmental stress (Wellman et al., 2007). Also in humans, the low-expressing polymorphic variant of the promoter region of the *HTT* gene has been associated with an elevated rate of posttraumatic stress disorders (H. J. Lee et al., 2005) and heightened risk of major depression in response to stressful life events (Caspi and Moffitt, 2006).

### Hypocretin controls sleep homeostasis in 5-HTT-deficient mice

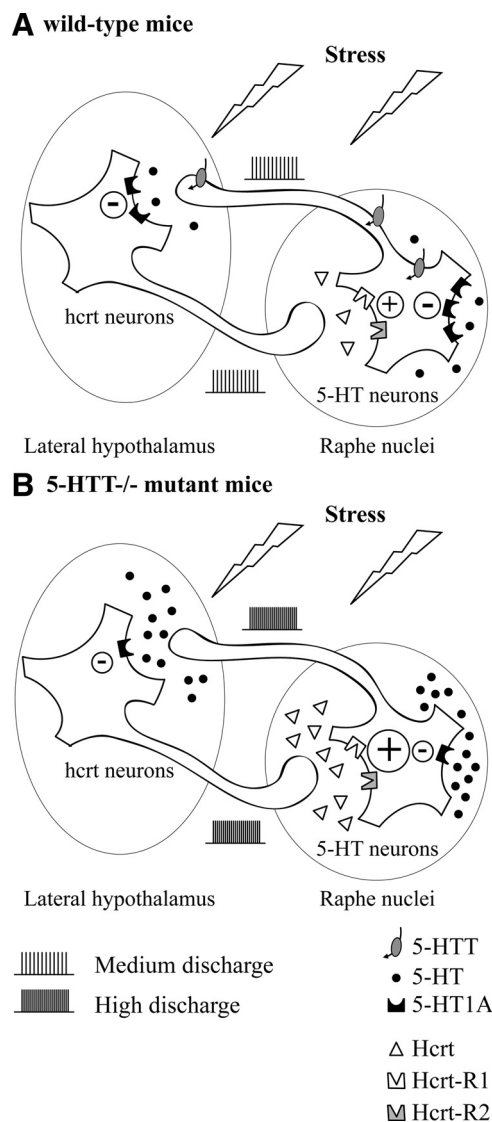
Numerous studies demonstrated that reciprocal influences exist between hypocretins and the HPA axis (Spinazzi et al., 2006). Administration of hcrt-1 activates the HPA axis (Hagan et al., 1999; Jászberényi et al., 2000; Kuru et al., 2000; Al-Barazani et al., 2001; Russell et al., 2001), and, conversely, stress increases preprohcrt mRNA contents in mice (Reyes et al., 2003) and activates hypocretin neurons in both rats and mice (Estabrooke et al., 2001; Espana et al., 2003; Winsky-Sommerer et al., 2004). In agreement with these data, we found here that RS activated hypocretinergic neurotransmission in both wild-type and *5-HTT*<sup>-/-</sup>



mice (as indicated by *c-Fos* labeling and *hcrt-R1* mRNA expression). However, this effect was stronger in mutant mice as reflected by the larger long-lasting influence of RS on hypocretin neuron activation and *hcrt-1* contents. Thus, at baseline, *5-HTT*<sup>-/-</sup> mutant mice exhibited in the anterior raphe area higher tissue levels of *hcrt-1* than wild-type mice, suggesting that 5-HT may affect *hcrt-1* production and/or degradation. After RS, these alterations were exacerbated in *5-HTT*<sup>-/-</sup> mice as indicated by the additional increase in *hcrt-1* tissue contents. However, the finding that hypocretinergic neurotransmission is enhanced in *5-HTT*<sup>-/-</sup> mice, notably after RS, is apparently contradictory with the well established inhibitory role of 5-HT on hypocretin neuron activity through 5-HT<sub>1A</sub> receptor activation (Muraki et al., 2004). Indeed, this discrepancy might be accounted for by the decreased density and function of 5-HT<sub>1A</sub> receptors in the hypothalamus of *5-HTT*<sup>-/-</sup> mice (Li et al., 1999, 2000; Alexandre et al., 2006), thereby causing a decreased inhibitory influence of 5-HT on hypocretin neurons. In turn, the hypocretinergic tone on 5-HT neurons would be enhanced, notably through elevated *hcrt-1* contents in the anterior raphe area of *5-HTT*<sup>-/-</sup> mice. Thus, our findings lead to the idea that the REMS altered response in *5-HTT*<sup>-/-</sup> mice might be related to an enhanced hypocretinergic neurotransmission in *5-HTT*<sup>-/-</sup> mice in response to stress.

This hypothesis is strongly supported by our observation that, although it had no impact on its own, blockade of *hcrt-R1* during stress in mutants rescued almost completely the delayed increase in REMS after stress, thereby mimicking the wild-type phenotype. This finding suggests that, in *5-HTT*<sup>-/-</sup> mice, the impaired REMS response to stress is mediated, at least in part, by hypocretin neurons via *hcrt-R1* receptors. This impairment seems specific to mutants because, in wild-type mice, acute blockade of *hcrt-R1* before stress did not modify the sleep response to RS. Thus, the adaptive regulation of sleep that follows an acute stress challenge requires a balance between the hypocretin and the 5-HT systems, which is impaired in *5-HTT*<sup>-/-</sup> mice. Here, we showed that *hcrt-R1* are involved in this balance, but this does not exclude a role for *hcrt-R2* also. Indeed, *hcrt-R2* are present on 5-HT neurons and may modulate the HPA axis during the stress response (Chang et al., 2007). This latter point needs to be addressed with selective *hcrt-R2* antagonists, but such ligands are not yet available. Finally, although other components of the 5-HT neurotransmission might be involved, directly or indirectly, in our paradigm, we propose a key role for 5-HT<sub>1A</sub> receptors in the deficit of REMS homeostasis after stress in *5-HTT*<sup>-/-</sup> mutants (Fig. 6). Indeed, (1) as observed in the latter mutants, knock-out mice lacking the 5-HT<sub>1A</sub> receptors exhibit no delayed REMS increase after RS (Boutrel et al., 2002), and (2) 5-HT<sub>1A</sub> receptors are downregulated and/or desensitized in the DR (Fabre et al., 2000; Bouali et al., 2003) and the hypothalamus (Li et al., 1999, 2000; Alexandre et al., 2006) of *5-HTT*<sup>-/-</sup> mutants. However, this model (Fig. 6) does not exclude a role for 5-HT<sub>1B</sub> receptors through regulation of 5-HT release at the terminal level, because 5-HT<sub>1B</sub> knock-out mice also exhibit a deficit in REMS homeostasis after RS (Adrien et al., 2004).

As mentioned above, in addition to sleep disorders, *5-HTT*<sup>-/-</sup> mice exhibit several behavioral alterations, including increased anxiety- and depression-like behaviors and vulnerability to stress (Hariri and Holmes, 2006). Conversely, mice with increased 5-HT reuptake exhibit a low anxiety-like phenotype (Jennings et al., 2006). Whether such behavioral changes may also be related to alterations in hypocretinergic neurotransmission is a relevant question to be addressed notably because several other animal models with sleep and/or behavioral alterations also exhibit



**Figure 6.** Model of *hcrt*–5-HT interactions in *5-HTT*<sup>-/-</sup> mutants and *5-HTT*<sup>+/+</sup> wild-type mice: influence of stress. **A**, In wild-type mice, restraint stress activates both *hcrt* and 5-HT neurons. Whereas *hcrt* excites 5-HT neurons through *hcrt-R1* and *hcrt-R2* receptors, 5-HT inhibits the activity of *hcrt* neurons through 5-HT<sub>1A</sub> receptors. In this model, the inhibitory influence of the indolamine on *hcrt* and 5-HT neurons (via 5-HT<sub>1A</sub> autoreceptors) will result in a new steady-state level and would allow the stress-enhanced sleep response (i.e., the delayed increase in REMS). **B**, In *5-HTT*<sup>-/-</sup> mice, hypocretinergic neurotransmission is enhanced at baseline. Such activation can result from adaptive changes in serotonergic neurotransmission and, notably, 5-HT<sub>1A</sub> receptor downregulation and/or desensitization in the hypothalamus that is triggered by the excess of extracellular 5-HT in mutant mice. In addition, these mice exhibit strong downregulation and desensitization of 5-HT<sub>1A</sub> autoreceptors in the dorsal raphe nucleus. We propose that this reduced 5-HT<sub>1A</sub>-mediated inhibitory influence of 5-HT on both 5-HT and *hcrt* neurons contributes to higher RS-induced enhancement of hypocretinergic and serotonergic neurotransmissions in *5-HTT*<sup>-/-</sup> versus wild-type mice. Such alterations of *hcrt*–5-HT interactions may account for the maladaptive REMS response to stress in *5-HTT*<sup>-/-</sup> mice (i.e., the absence of the delayed increase in REMS), because pharmacological blockade of *hcrt-R1* restored the delayed increase in REMS normally observed after restraint stress.

changes in the hypocretinergic system (Taheri et al., 2001; Allard et al., 2004; Feng et al., 2007; Lutter et al., 2008). Interestingly, studies in humans showed that *hcrt-1* levels in the CSF (1) negatively correlate with symptoms such as lassitude in suicide attempters (Brundin et al., 2007), (2) are slightly increased in depressed patients, and (3) are reduced by treatment with selective serotonin reuptake inhibitors (Salomon et al., 2003). Thus, as

suggested in the present study, hypocretin neurons, which are under the influence of 5-HT, may modulate the emotional state.

## Conclusion

Our study using 5-HTT<sup>-/-</sup> mice is reminiscent of human pathology in which low functioning of the 5-HTT is associated with increased vulnerability to stress and heightened risk of developing depression. Accordingly, the hypocretin/5-HT interactions would be involved not only in sleep regulation but also in the control of mood- and emotion-driven functions and may underlie psychoaffective disorders.

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