

Optogenetic Manipulation of Activity and Temporally Controlled Cell-Specific Ablation Reveal a Role for MCH Neurons in Sleep/Wake Regulation

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Melanin-concentrating hormone (MCH) is a neuropeptide produced in neurons sparsely distributed in the lateral hypothalamic area. Recent studies have reported that MCH neurons are active during rapid eye movement (REM) sleep, but their physiological role in the regulation of sleep/wakefulness is not fully understood. To determine the physiological role of MCH neurons, newly developed transgenic mouse strains that enable manipulation of the activity and fate of MCH neurons *in vivo* were generated using the recently developed knockin-mediated enhanced gene expression by improved tetracycline-controlled gene induction system. The activity of these cells was controlled by optogenetics by expressing channelrhodopsin2 (E123T/T159C) or archaerhodopsin-T in MCH neurons. Acute optogenetic activation of MCH neurons at 10 Hz induced transitions from non-REM (NREM) to REM sleep and increased REM sleep time in conjunction with decreased NREM sleep. Activation of MCH neurons while mice were in NREM sleep induced REM sleep, but activation during wakefulness was ineffective. Acute optogenetic silencing of MCH neurons using archaerhodopsin-T had no effect on any vigilance states. Temporally controlled ablation of MCH neurons by cell-specific expression of diphtheria toxin A increased wakefulness and decreased NREM sleep duration without affecting REM sleep. Together, these results indicate that acute activation of MCH neurons is sufficient, but not necessary, to trigger the transition from NREM to REM sleep and that MCH neurons also play a role in the initiation and maintenance of NREM sleep.

Key words: ablation; cell fate; channelrhodopsin2; hypothalamus; optogenetics; REM sleep

Introduction

Melanin-concentrating hormone (MCH)-producing neurons are coexpressed with orexin/hypocretin-producing neurons in

the lateral hypothalamic area (LHA) but are more numerous and extend further rostrocaudally (Bittencourt et al., 1992; Peyron et al., 1998; Elias et al., 2008). MCH neurons project widely throughout the brain and densely innervate the cholinergic and monoaminergic arousal centers (Bittencourt et al., 1992). MCH neurons have been reported to be GABAergic (Del Cid-Pellitero and Jones, 2012). While MCH receptor-1 (MCHR1; the only receptor found in rodents) binding activates G_q, G_i, and G_o subunits (Hawes et al., 2000), the major effect of MCHR1 binding is to decrease cAMP levels (Chambers et al., 1999; Lembo et al., 1999) and cellular electrophysiological studies have revealed presynaptic and postsynaptic inhibitory effects of MCH (Gao and van den Pol, 2001; Wu et al., 2009). Thus, MCH neurons exert neuroinhibitory effects on downstream targets.

MCH has been implicated in several functions, including feeding, energy balance, and locomotor activity (Shimada et al., 1998; Asakawa et al., 2002; Marsh et al., 2002; Semjonous et al., 2009); its involvement in sleep and wakefulness has also been extensively studied. Intracerebroventricular injections of MCH early in the dark period increased rapid eye movement (REM) sleep in a dose-dependent manner (Verret et al., 2003); some

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doses also increased non-REM (NREM) sleep. A recent study reported that optogenetic stimulation of MCH neurons increased both NREM and REM sleep (Konadhode et al., 2013). These results are consistent with the observation that intracerebroventricular injections of MCH increase both sleep states, but are at variance with results from functional neuroanatomical, pharmacological (Ahnaou et al., 2008), and optogenetic (Jego et al., 2013) studies that specifically indicate a role for MCH neurons in REM sleep regulation.

To this point, the precise role of MCH neurons in REM sleep regulation remains unclear and the involvement of MCH neurons in NREM sleep is uncertain (Jego and Adamantidis, 2013; Jones and Hassani, 2013; Luppi et al., 2013b; McGinty and Alam, 2013; Pelluru et al., 2013). One potential source of the discrepant results from the optogenetic studies could be the number of MCH neurons transfected with the light-sensitive protein and subsequently stimulated. Therefore, we generated new transgenic mice (Berndt et al., 2011; Han et al., 2011; Tanaka et al., 2012) to control MCH neurons and bred this strain with three other transgenic lines to create novel bigenic strains that enabled cell-specific manipulation of MCH neurons: an E123T/T159C double mutant of channelrhodopsin2 (ChR2) for stimulation (Berndt et al., 2011), archaerhodopsinTP009 (ArchT) for inhibition (Han et al., 2011), and diphtheria toxin A (DTA) for ablation. We find that optogenetic stimulation of MCH neurons induced REM sleep with a short latency and that continuous stimulation resulted in a net increase of REM sleep in conjunction with a decrease of NREM sleep. Although optogenetic inhibition of MCH neurons had no effect on sleep/wakefulness, conditional ablation of MCH neurons resulted in hyposomnia. These results demonstrate a critical role for MCH neurons in REM sleep regulation under acute stimulation and provide unequivocal evidence for this LHA neuronal population in sleep/wake control.

Materials and Methods

Animal usage. All experimental procedures involving animals were approved by the animal care and use committees at Nagoya University, Hokkaido University, and SRI International and were in accordance with National Institutes of Health guidelines. All efforts were made to minimize animal suffering or discomfort and to reduce the number of animals used.

Generation of MCH-tetracycline-controlled transactivator bacterial artificial chromosome transgenic mice. The codons of bacterial tetracycline repressor protein and viral VP16 activator domain [tetracycline-controlled transactivator (tTA)] were fully mammalianized (Inamura et al., 2012). Mouse bacterial artificial chromosome (BAC) DNA (clone RP23-348H6) was initially modified by inserting an RpsI-Zeo cassette (a gift from Dr. Hisashi Mori, Toyama University) into the translation initiation site of the MCH gene followed by the replacement with a cassette containing tTA and the SV40 polyadenylation signal. BAC DNA was linearized by P1-SceI enzyme digestion (New England Biolabs) and injected into fertilized eggs from BDF1 mice. Two founders were obtained and line B was used in this report.

Generation of tetracycline operator ChR2 (E123T/T159C) knock-in mice. To achieve sufficient opsin expression (ChR2) in MCH neurons to drive photocurrents, we used a knockin-mediated enhanced gene expression by improved tetracycline-controlled gene induction (KENGE-tet) strategy (Tanaka et al., 2012). We used the same housekeeping gene, *actb*, but the position of gene targeting was 800 bp downstream of the polyA signal, which was different from our initial report (Tanaka et al., 2012).

We constructed a plasmid containing the tetracycline operator (TetO) ChR2 polyA cassette with the Neo selection gene flanked on both sides by flippase recombinase target (FRT) sites in which the following elements were connected in tandem: TetO sequence, rabbit β -globin intron, ChR2(E123T/T159C) enhanced yellow fluorescent protein (EYFP) cDNA, SV40 polyadenylation signal, and FRT-flanked phosphoglycerate

kinase-EM7-Neo. To insert the above cassette downstream of the mouse β -actin gene, a 265 bp 5' homology arm consisting of the sequence from 535 to 799 bp downstream of the mouse β -actin gene polyadenylation signal (AATAAA) was connected to the 5' end of the TetO sequence, and a 265 bp 3' homology arm consisting of the sequence from 800 to 1064 bp downstream of the AATAAA sequence was connected to the 3' end of Neo. To perform BAC recombination, DNA fragments with both homology arms were electroporated into bacteria carrying the BAC (clone RP23-289L7) and the pBADTcTypeG plasmid (a gift from Dr. Manabu Nakayama, Kazusa Institute, Japan; Nakayama and Ohara, 2005). Kanamycin-resistant clones were selected as the modified BAC and the TetO ChR2 polyA cassette was subsequently inserted 800 bp downstream of mouse β -actin gene AATAAA. The targeting vector was isolated from the kanamycin-resistant, modified BAC clone using a retrieval technique involving insertion into the pMCS-DTA plasmid (a gift from Dr. Kosuke Yusa, Osaka University, Japan).

We subsequently obtained a targeting vector comprising a 11 kb 5' homology arm, TetO ChR2 polyA cassette with Neo, a 1.5 kb 3' homology arm, and a DTA subunit. EGR-G01 ES cell (established by F1 from 129S2 and C57BL/6-Tg(CAG/Acr-EGFP)) cells were used. We obtained 28 recombinant clones out of 96 G418-resistant clones. Recombination was confirmed by Southern blotting with a 424 bp 3-prime outside probe, which recognized an 8.1 kb fragment of the wild-type allele and a 6 kb fragment of the targeted allele in genomic DNA digested with XbaI. Germline-transmitted offspring were established as TetO ChR2-Neo knock-in mice. TetO ChR2-Neo mice were crossed with ROSA-Flp mice (Farley et al., 2000), and FRT-flanked Neo selection markers were removed. *TetO ChR2* knock-in mice were subsequently generated.

Generation of bigenic MCH mouse strains. To express ChR2, ArchT, or DTA in MCH neurons, *MCH-tTA* mice were bred with *TetO ChR2* mice, *TetO ArchT* mice (Tsunematsu et al., 2013), or *TetO DTA* mice (B6.Cg-Tg(tetO-DTA)1Gfi/J, 008468, Jackson Laboratory), respectively.

Brain slice preparation. Male and female *MCH-tTA*; *TetO ChR2* or *MCH-tTA*; *TetO ArchT* bigenic mice (3–6 weeks old) were used for whole-cell recordings. The mice were deeply anesthetized with isoflurane (Abbott Japan) and decapitated. Brains were quickly isolated in ice-cold cutting solution consisting of (in mM) the following: 280 sucrose, 2 KCl, 10 HEPES, 0.5 CaCl₂, 10 MgCl₂, 10 glucose, pH 7.4 with NaOH, bubbled with 100% O₂. Brains were cut coronally into 350 μ m slices with a microtome (VTA-1200S, Leica). Slices containing the LHA were transferred to an incubation chamber shielded from light and filled with a physiological solution containing the following (in mM): 135 NaCl, 5 KCl, 1 CaCl₂, 1 MgCl₂, 10 HEPES, 10 glucose, pH 7.4 with NaOH, bubbled with 100% O₂. This was incubated for \geq 1 h at room temperature (RT; 24–26°C).

In vitro electrophysiological recordings. At RT, the slices were transferred to a recording chamber (RC-27L, Warner Instrument) on a fluorescence microscope stage (BX51WI, Olympus). Neurons having EGFP and EYFP fluorescence were identified as MCH neurons and were subjected to electrophysiological recordings. The fluorescence microscope was equipped with an infrared camera (C2741-79, Hamamatsu Photonics) for infrared differential interference contrast imaging and a cooled charge-coupled device (CCD) camera (Cascade 650, Roper Scientific) for fluorescent imaging. Each image was displayed separately on a monitor and saved on a computer. Recordings were performed with an Axopatch 200B amplifier (Molecular Devices) using a borosilicate pipette (GC150-10, Harvard Apparatus) prepared by a micropipette puller (P-97, Sutter Instruments), and filled with intracellular solution (4–6 M Ω) consisting of (in mM) the following: 138 K-gluconate, 10 HEPES, 8 NaCl, 0.2 EGTA-Na₃, 2 MgATP, 0.5 Na₂GTP, pH 7.3 with KOH. The osmolality of the solution was checked by a vapor pressure osmometer (model 5520, Wescor). The osmolality of the internal and external solutions was 280–290 and 320–330 mOsm/l, respectively. The liquid junction potential of the patch pipette and perfused extracellular solution was estimated to be 16 mV and was corrected in the data. Recording pipettes were under positive pressure while advancing toward individual cells in the slice, and tight seals on the order of 1.0–1.5 G Ω were made by negative pressure. The membrane patch was then ruptured by suction. The series resistance during recording was 10–25 M Ω . The reference electrode was an Ag-AgCl pellet immersed in bath solution. During recordings at RT, cells

were superfused with extracellular solution at a rate of 1.6 ml/min using a peristaltic pump (Dynamax, Rainin). Blue (475 ± 17.5 nm, 2.5 mW) and green light (549 ± 7.5 nm, 0.9 mW) was generated by a SPECTRA light engine (Lumencor).

The output signal was low-pass filtered at 5 kHz and digitized at 10 kHz. Data were recorded on a computer through a Digidata 1322A analog-to-digital converter using pClamp software version 10.2 (Molecular Devices).

Drugs. Tetrodotoxin (TTX; Wako) was dissolved in extracellular solution at a concentration of $1 \mu\text{M}$. During experiments, TTX was applied by bath application.

Immunohistochemistry. Male and female *MCH-tTA*; *TetO ChR2* and *MCH-tTA*; *TetO ArchT* bigenic mice (6 weeks old) were deeply anesthetized with isoflurane and perfused sequentially with 20 ml of chilled saline and 20 ml of chilled 10% formalin solution (Wako). The brains were removed and immersed in the above fixative solution for 24 h at 4°C and then immersed in a 30% sucrose solution for ≥ 2 d. The brains were quickly frozen in embedding solution (Sakura Finetech). For MCH and GFP double staining, coronal sections ($40 \mu\text{m}$) of *MCH-tTA*; *TetO ChR2* and *MCH-tTA*; *TetO ArchT* bigenic mice brains were incubated with mouse anti-GFP antiserum (1:1000; Wako) for 24 h at 4°C . These sections were incubated with Alexa 488-labeled donkey anti-mouse IgG (1:1000; Biotium) for 1 h at RT. The sections were then incubated with rabbit anti-MCH antiserum (1:2000; Sigma-Aldrich) for 24 h at 4°C and incubated with Alexa 594-labeled donkey anti-rabbit IgG (1:1000; Biotium) for 1 h at RT. To confirm the specificity of antibodies, incubations without primary antibody were conducted as a negative control in each experiment and, in each case, no signal was observed.

To determine the number of MCH and orexin neurons, coronal sections ($40 \mu\text{m}$) were stained by the avidin-biotin-peroxidase method. Brain sections were incubated for 40 min in phosphate buffer containing 0.3% H_2O_2 to inactivate endogenous peroxidase. Sections were transferred into PBS containing 0.25% Triton X-100 and 1% bovine serum albumin fraction V (PBS-BX) for 30 min and then incubated with rabbit anti-MCH antiserum (Sigma-Aldrich) diluted 1:4000 in PBS-BX, goat anti-orexin IgG antibody (Santa Cruz Biotechnology) diluted 1:2000 in PBS-BX overnight at 4°C . Sections were then incubated with biotin-labeled goat anti-rabbit IgG antibody (1:1000; Vector Laboratories) or biotin-labeled horse anti-goat IgG antibody (1:1000; Vector Laboratories) for 1 h at RT followed by incubation with avidin and biotinylated peroxidase complex solution for 1 h at RT. Bound peroxidase was visualized by DAB-buffer tablet (Merck) with 0.0015% H_2O_2 , resulting in a golden-brown reaction product. Approximately 10 slices from a one-in-four series were selected for counting of MCH and orexin neurons. Sections were mounted and examined with a fluorescence microscope (BZ-9000, Keyence) or a confocal microscope (LSM710, Zeiss). Cell counts were made bilaterally. The number of MCH neurons in the brain is presented as the sum of the number of MCH neurons counted in every fourth brain slice at a thickness of $40 \mu\text{m}$.

In vivo photo illumination using freely moving mice. Male *MCH-tTA*; *TetO ChR2* and *MCH-tTA*; *TetO ArchT* bigenic mice, 12 weeks of age or older, were housed under controlled lighting (12 h light/dark cycle; lights on from 8:00 to 20:00) and temperature ($22 \pm 3^\circ\text{C}$) conditions. Food and water were available *ad libitum*. Mice were anesthetized with isoflurane using a vaporizer for small animals (Bio Research Center) and positioned in a stereotaxic frame (David Kopf Instruments). Plastic fiber optics (0.5 mm in diameter; Eska, Mitsubishi Rayon) were bilaterally implanted into the hypothalamus ~ 1 mm above the LHA (1.2 mm posterior, 1 mm lateral from bregma, 3.5 mm depth from brain surface). Electrodes for EEG and EMG were implanted on the skull and neck muscles, respectively. For the head-fixed mice, a U-shaped plastic plate was attached to the skull to enable fixation of the mouse's head to the stereotaxic frame during recordings. These were attached to the skull using dental cement (GC). The mice were then housed separately for a recovery period of ≥ 7 d.

Continuous EEG and EMG recordings were performed through a slip ring (Air Precision, Le Pressis Robinson) designed so that the movement of the mouse was unrestricted. EEG and EMG signals were amplified (AB-610J, Nihon Koden), filtered (EEG, 1.5–30 Hz; EMG, 15–300 Hz), digitized at a sampling rate of 128 Hz, and recorded using SleepSign

software version 3 (Kissei Comtec). Blue (475 ± 17.5 nm) and green light (542 ± 13.5 nm) was generated by the SPECTRA light engine and applied through plastic optical fibers bilaterally inserted 1 mm above the LHA. An optical swivel (COME2, Lucir) was used for unrestricted *in vivo* photo illumination. The power intensities of blue and green light at the tip of the plastic fiber optics (0.5 mm diameter) were 25.8 and 33.4 mW/mm^2 , respectively, as measured by a power meter (VEGA, Ophir Optonics). Each animal's behavior was monitored through a CCD video camera and recorded on a computer synchronized with EEG and EMG recordings using the SleepSign video option system (Kissei Comtec). The infrared activity monitor was a sensor mounted on top of the cage to measure locomotor activity. The sensor's output signals (representing the magnitude of each animal's movement) were digitally converted and transferred to a computer.

MCH-tTA; TetO DTA bigenic mouse surgery. *MCH-tTA*; *TetO DTA* bigenic mice were housed under controlled lighting (12 h light/dark cycle; lights on from 7:00 to 19:00) and temperature (22°C) conditions. Food and water were available *ad libitum*. Doxycycline-containing chow (Dox chow) was made by adding 10% doxycycline (Dox) powder (Kyoritsu Seiyaku) to normal chow (Labo MR Stock, Nosan) at a final concentration of 100 mg/kg. Labo MR stock was provided during the Dox(–) period (from 10 weeks of age). Mating pairs of *MCH-tTA* mice and *TetO DTA* mice were fed with Dox-containing chow [Dox(+) condition] from the day of mating. During the prenatal and early postnatal periods, Dox was supplied via maternal circulation or lactation, respectively. After weaning, *MCH-tTA*; *TetO DTA* mice were fed with Dox(+) chow until the day of the experiment.

MCH-tTA; TetO DTA bigenic mice (8 weeks of age) were anesthetized with an intraperitoneal injection of a drug mixture containing ketamine (75 mg/kg) and medetomidine (1 mg/kg), placed in a stereotaxic device, and chronically implanted with EEG and EMG electrodes for polysomnographic recording of sleep and wakefulness states. Two stainless-steel screws (diameter, 1.0 mm) were implanted into the skull over the left cerebral hemisphere (frontal: 1.5 mm lateral to midline, 1.0 mm anterior to bregma; parietal: 1.5 mm lateral to midline, 1.0 mm anterior to lambda) according to the atlas of Franklin and Paxinos (1997). A third screw was placed over the frontal bone to serve as a ground electrode. EMG activity was monitored using stainless-steel Teflon-coated wires inserted bilaterally into the neck muscles. All electrodes were attached to a microconnector, and the entire assembly was fixed to the skull with dental cement. Mice were injected with indomethacin (1 mg/kg, i.p.) and penicillin G (200 U/kg, i.m.) immediately after surgery. All procedures were performed on a heating pad, and mice were allowed to recover for 10–14 d before being transferred to sleep recording chambers.

Vigilance state determination. Polysomnographic recordings were automatically scored offline as wakefulness, NREM sleep, or REM sleep by SleepSign, in 4 s epochs (optogenetics experiments) or 10 s epochs (ablation experiments), according to standard criteria (Tobler et al., 1997; Yamanaka et al., 2002). All vigilance state classifications assigned by SleepSign were examined visually and corrected if necessary. The same individual, blinded to genotype and experimental condition, scored all EEG/EMG recordings. Spectral analysis of the EEG was performed by fast Fourier transform (FFT; sampled at 128 Hz). This analysis yielded a power spectral profile over a 0–25 Hz window with a 1 Hz resolution divided into delta (1–5 Hz), theta (6–10 Hz), alpha (10–13 Hz), and beta (13–25 Hz) waves.

Statistical analysis. Data were analyzed by paired *t* test, unpaired *t* test, one-way ANOVA, two-way ANOVA, or nonparametric Kruskal–Wallis test (as appropriate for the parameters examined) using KaleidaGraph 4.0 software (Hulinks). When appropriate, ANOVA tests were followed by *post hoc* analysis of significance using Fisher's protected least significant difference test or Bonferroni's test. *p* values < 0.05 were considered statistically significant.

Results

Specific activation of MCH neurons using ChR2

MCH neurons in the brains of *MCH-tTA*; *TetO ChR2* bigenic mice specifically expressed ChR2 (Fig. 1A) as confirmed by double-labeled immunohistochemistry. An anti-GFP antibody

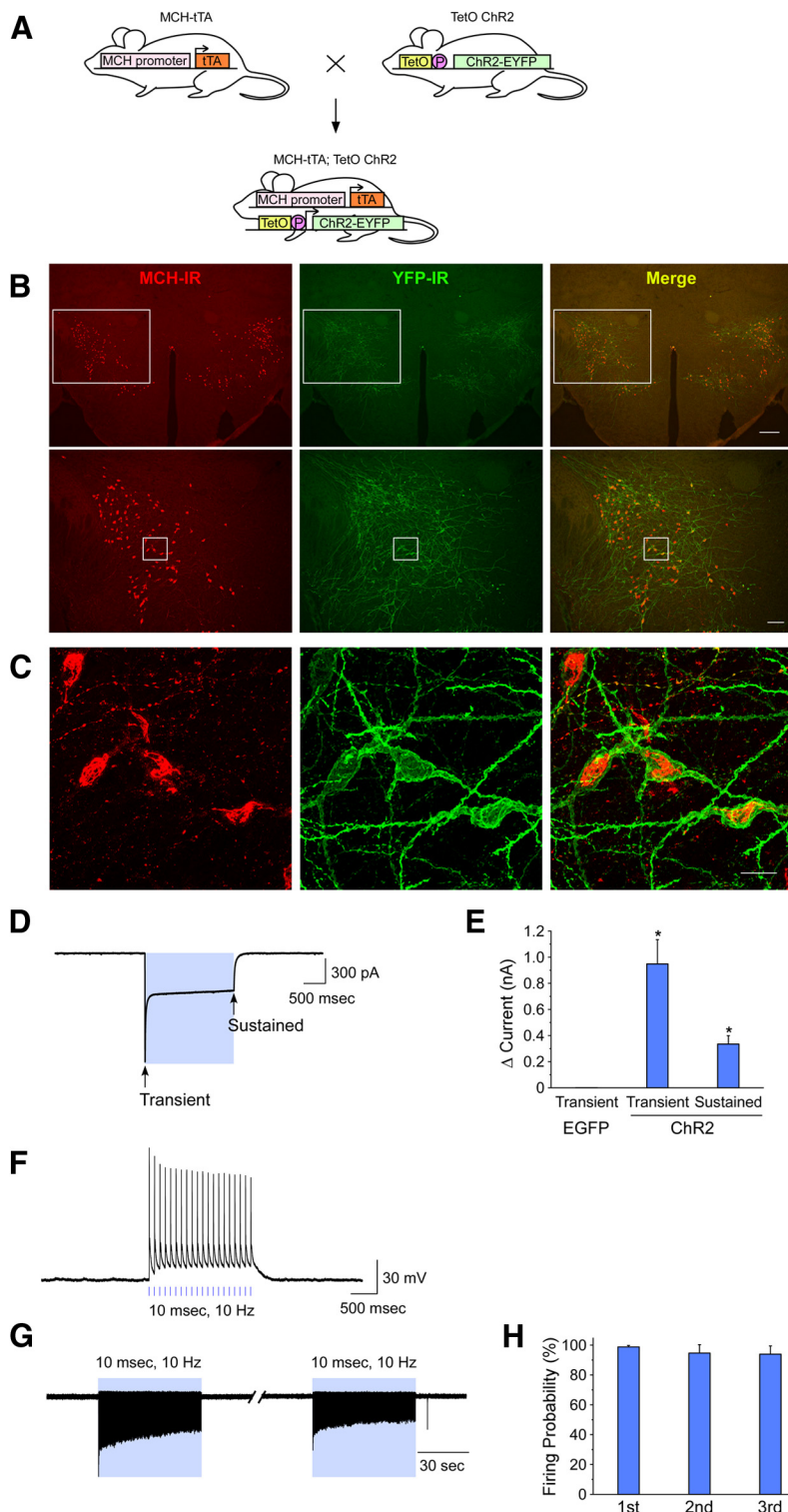


Figure 1. Specific activation of MCH neurons. **A**, Schematic showing generation of bigenic *MCH-tTA*; *TetO ChR2* mice. P, Minimal promoter. **B**, **C**, Immunohistochemical analyses revealed that ChR2 is specifically expressed in MCH neurons in the *MCH-tTA*; *TetO ChR2* bigenic mouse brain. **B**, MCH-immunoreactive (MCH-IR) neurons (left, Alexa594, red) and ChR2::EYFP-immunoreactive (YFP-IR) neurons (middle, Alexa488, green) located in the LHA and the zona incerta. Right, Merged image shows specific expression of ChR2 in MCH neurons. Bottom row represents higher magnifications of the regions enclosed by the squares in the top row. Scale bars: (in top, right) top row, 200 μm ; (in bottom, right) bottom row, 100 μm . **C**, Confocal microscopic image of the region indicated by the squares in **B**, bottom row. Scale bar, 20 μm . **D–H**, Slice patch-clamp recordings from MCH neurons. **D**, Photocurrent induced by blue light (475 ± 17.5 nm, 2.5 mW, 2 s; $V_h = -60$ mV, TTX 1 μM). **E**, Bar graph summarizing the data obtained from **D**. **F**, Current-clamp mode recording, blue light pulses (2.5 mW, 10 ms, 10 Hz) generated action potentials in MCH neurons. **G**, Loose cell-attached recording from MCH neurons. Blue light pulses (2.5 mW, 10 ms, 10 Hz) for 1 min every 5 min generated action potentials. **H**, Summary data from **G**. Values represent means \pm SEM. * $p < 0.05$ versus EGFP-expressing MCH neurons.

was used to detect ChR2-EYFP fusion proteins. Merged pictures show that ChR2-EYFP was exclusively observed in MCH neurons in bigenic *MCH-tTA*; *TetO ChR2* mice (Fig. 1*B,C*). Ectopic expression of ChR2 outside of MCH neurons was not observed. Confocal microscopic observation revealed that ChR2 was located in the plasma membrane of somata and dendrites of MCH neurons. MCH neurons expressing ChR2 did not show blebbing or other features indicative of inappropriate trafficking (Fig. 1*C*). The number and morphology of MCH neurons in *MCH-tTA*; *TetO ChR2* mice were indistinguishable from those of monogenic littermate mice (*MCH-tTA* or *TetO ChR2* mice, data not shown), suggesting that ChR2 expression is not toxic to MCH neurons. The ChR2 expression rate (YFP-immunoreactive/MCH-immunoreactive $\times 100\%$) in *MCH-tTA*; *TetO ChR2* mice was $88.0 \pm 1.3\%$ ($n = 9$).

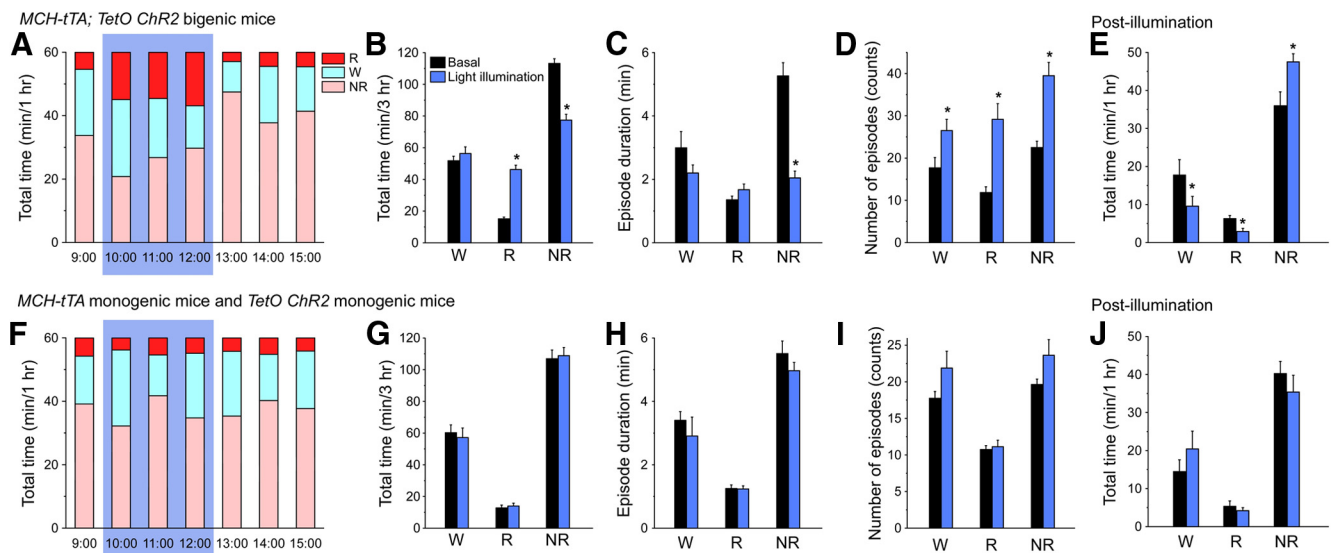
To confirm the function of ChR2 in MCH neurons, slice patch-clamp analyses were performed. Most MCH neurons did not exhibit spontaneous firing (0.0 ± 0.0 Hz, $n = 5$). Under whole-cell voltage-clamp mode, blue light (475 ± 17.5 nm, 2.5 mW) was illuminated through the objective lens in the presence of TTX. At a holding potential of -60 mV, blue light illumination for 2 s induced robust inward currents (Fig. 1*D*). Blue light illumination-induced transient and sustained currents were 948.2 ± 185.5 pA ($n = 9$, $p < 0.001$, ANOVA) and 335.0 ± 64.3 pA ($n = 9$, $p = 0.04$, ANOVA), respectively (Fig. 1*E*). Under current-clamp mode, a blue light pulse (10 ms, 10 Hz) was applied. Blue light illumination instantaneously caused depolarization and action potentials were generated in conjunction with light pulses (Fig. 1*F*). Next, activation of MCH neurons was confirmed using loose cell-attached recordings, which recorded firing frequencies without affecting intracellular conditions. Blue light illumination for 1 min (10 ms, 10 Hz) evoked repetitively induced action potentials with high fidelity (Fig. 1*G,H*). These results strongly suggested that blue light illumination would activate MCH neurons *in vivo*.

Activation of MCH neurons increases total time in REM sleep

To study the physiological significance of MCH neuronal activity in the regulation of sleep/wakefulness, MCH neurons were activated *in vivo* using freely moving *MCH-tTA*; *TetO ChR2* mice. To determine sleep/wakefulness states, mice were

Table 1. Vigilance state parameters recorded from *MCH-tTA; TetO Chr2* bigenic mice, monogenic littermate mice, and *MCH-tTA; TetO ArchT* bigenic mice under basal conditions

| | Wakefulness | | | REM sleep | | | NREM sleep | | |
|-----------------------------|---------------|---------------------------|----------------------------|------------|---------------------------|----------------------------|--------------|---------------------------|----------------------------|
| | Control | <i>MCH-tTA; TetO Chr2</i> | <i>MCH-tTA; TetO ArchT</i> | Control | <i>MCH-tTA; TetO Chr2</i> | <i>MCH-tTA; TetO ArchT</i> | Control | <i>MCH-tTA; TetO Chr2</i> | <i>MCH-tTA; TetO ArchT</i> |
| 24 h | | | | | | | | | |
| Total time (min) | 792.5 ± 40.7 | 734.5 ± 30.3 | 755.9 ± 14.8 | 61.5 ± 6.8 | 81.2 ± 9.3 | 80.8 ± 10.6 | 586.0 ± 10.6 | 624.2 ± 25.1 | 603.2 ± 12.3 |
| Episode duration (s) | 479.1 ± 91.7 | 479.3 ± 81.9 | 546.7 ± 86.9 | 65.8 ± 4.8 | 75.1 ± 1.6 | 71.7 ± 4.7 | 281.6 ± 26.2 | 251.6 ± 12.1 | 251.2 ± 13.2 |
| Number of episodes (counts) | 118.0 ± 4.6 | 108.1 ± 8.1 | 131.4 ± 11.4 | 52.6 ± 3.4 | 48.7 ± 4.6 | 59.4 ± 6.6 | 133.3 ± 5.1 | 127.0 ± 6.8 | 148.2 ± 10.5 |
| Light period | | | | | | | | | |
| Total time (min) | 232.4 ± 14.7 | 210.2 ± 9.2 | 204.9 ± 5.9 | 50.5 ± 8.1 | 62.5 ± 5.0 | 68.1 ± 8.1 | 437.1 ± 16.5 | 447.3 ± 12.2 | 446.9 ± 6.2 |
| Episode duration (s) | 212.3 ± 25.9 | 167.3 ± 13.7 | 169.8 ± 30.7 | 71.5 ± 7.6 | 83.7 ± 3.7 | 79.8 ± 3.1 | 344.8 ± 42.8 | 285.7 ± 33.9 | 269.0 ± 15.4 |
| Number of episodes (counts) | 79.4 ± 9.0 | 80.9 ± 10.4 | 95.2 ± 5.7 | 39.5 ± 4.3 | 41.3 ± 3.8 | 48.2 ± 5.1 | 90.4 ± 9.8 | 97.7 ± 8.9 | 108.6 ± 6.1 |
| Dark period | | | | | | | | | |
| Total time (min) | 551.0 ± 43.9 | 524.4 ± 37.6 | 551.0 ± 16.8 | 12.7 ± 2.9 | 18.7 ± 5.7 | 12.7 ± 3.4 | 156.4 ± 41.3 | 176.9 ± 32.2 | 156.3 ± 14.9 |
| Episode duration (s) | 745.8 ± 204.8 | 791.3 ± 153.4 | 923.6 ± 147.3 | 60.2 ± 3.4 | 66.4 ± 2.9 | 63.6 ± 9.7 | 218.3 ± 12.6 | 217.4 ± 22.4 | 233.4 ± 24.6 |
| Number of episodes (counts) | 38.6 ± 8.2 | 27.3 ± 5.9 | 36.2 ± 7.2 | 13.1 ± 3.0 | 7.4 ± 2.0 | 11.2 ± 1.7 | 42.9 ± 9.4 | 29.3 ± 6.5 | 39.6 ± 6.4 |

**Figure 2.** *In vivo* optogenetic activation of MCH neurons increased time spent in REM sleep. **A, F**, Bar graphs illustrating the time in each vigilance state during the light period (9:00–16:00) in (**A**) *MCH-tTA; TetO Chr2* bigenic mice and (**F**) *MCH-tTA* monogenic mice and *TetO Chr2* monogenic mice. Blue light pulses (475 ± 17.5 nm, 25.8 mW/mm², 10 ms, 10 Hz for 3 h from 10:00 to 13:00) were applied. Blue background indicates the period of illumination. **B, G**, Bar graphs summarizing the data from **A** and **F**. The time spent in each vigilance state during illumination for 3 h is summarized. Black bars, No light illumination; blue bars, blue light pulse illumination. **C, H**, Mean episode duration. **D, I**, Number of episodes. **E, J**, Bar graph summarizing the time in each vigilance state for 1 h after cessation of illumination (from 13:00 to 14:00) in *MCH-tTA; TetO Chr2* bigenic mice (**E**) and *MCH-tTA* monogenic mice and *TetO Chr2* monogenic mice (**J**). W, Wakefulness; R, REM sleep; NR, NREM sleep. Values are represented as means \pm SEM. * $p < 0.05$ versus basal (no illumination).

chronically implanted with EEG and EMG electrodes and bilateral optical fibers. The tip of the optical fiber was stereotaxically placed 1 mm above MCH neurons to illuminate Chr2-expressing MCH neurons.

First, the vigilance state distribution of *MCH-tTA; TetO Chr2* bigenic mice under basal conditions was compared with that of littermate monogenic mice (*MCH-tTA* mice or *TetO Chr2* mice). There were no significant differences between these strains in the total time, mean episode duration, or the number of episodes of wakefulness, REM sleep, or NREM sleep (Table 1), indicating that Chr2 expression in MCH neurons had no effect on sleep/wakefulness patterns.

Blue light pulses (475 ± 17.5 nm, 25.8 mW/mm², 10 ms, 10 Hz) were applied during the first half of the light period (from 10:00 to 13:00) and *MCH-tTA; TetO Chr2* mice without light illumination were used as a comparison. The total time in REM sleep was significantly increased in *MCH-tTA; TetO Chr2* mice during activation of MCH neurons (Fig. 2*A, B*). During this 3 h period, the total time spent in wakefulness, REM sleep, and

NREM sleep under basal conditions and after activation of MCH neurons was 51.8 ± 2.9 min ($n = 6$) and 56.3 ± 4.2 min [$n = 6$, $p = 0.25$, not significantly different (NS) paired t test]; 15.1 ± 1.2 min ($n = 6$) and 46.3 ± 2.7 min ($n = 6$, $p < 0.001$, paired t test); and 113.1 ± 3.0 min ($n = 6$) and 77.4 ± 3.6 min (NREM, $n = 6$, $p < 0.001$, paired t test), respectively. These results indicated that activation of MCH neuronal activity resulted in increased time spent in REM sleep accompanied by a decrease in time spent in NREM sleep, but without affecting the time spent in wakefulness. The mean episode duration of NREM sleep was significantly decreased from 5.3 ± 0.4 min ($n = 6$) to 2.0 ± 0.2 min ($n = 6$, $p < 0.001$, paired t test) by activation of MCH neurons (Fig. 2*C*). Meanwhile, the number of episodes of wakefulness, REM sleep, and NREM sleep was significantly increased (Fig. 2*D*). These results suggest that activation of MCH neurons increased the probability of transitions from NREM to REM sleep, resulting in increased REM and decreased NREM sleep over the 3 h period. Since the duration of REM episodes was unchanged, however, an overall fragmentation of sleep/wakefulness resulted, as indicated

by the increased number of episodes of all states. After stimulation of MCH neurons ceased, NREM sleep was significantly increased with decreases in wakefulness and REM sleep (Fig. 2E), suggesting a compensatory rebound in NREM sleep.

Blue light illumination (475 ± 17.5 nm, 25.8 mW/mm², 10 ms, 10 Hz) had no effect on sleep/wakefulness in monogenic *MCH-tTA* or *TetO Chr2* mice (Fig. 2F). Neither the total time (Fig. 2G), nor mean episode duration (Fig. 2H), nor the number of episodes (Fig. 2I) during illumination for 3 h, nor the total time after illumination (Fig. 2J) of each vigilance state were significantly affected by blue light pulse.

Does activation of MCH neurons induce physiological REM sleep?

Although activation of MCH neurons increased the time spent in REM sleep as determined from EEG and EMG recordings in freely moving mice, it was unclear whether this state met criteria for physiological REM sleep other than those based on EEG and EMG recordings. To evaluate this further, eye and whisker movements during activation of MCH neurons were visually analyzed. To monitor eye and whisker movements during MCH neuron stimulation, mice were affixed to a stereotaxic frame via a plastic U-shape frame fixed onto the skull. After a week adaptation and training to these conditions, mice spontaneously cycle between wakefulness and sleep. While a mouse was in spontaneous NREM sleep, MCH neurons were activated and the EEG and EMG indicated signs of REM sleep shortly after initiation of stimulation (475 ± 17.5 nm, 25.8 mW/mm², 10 ms, 10 Hz). During MCH neuron activation-induced REM sleep, the eyes and whiskers moved intermittently, similar to that observed during spontaneously occurring REM sleep. These results suggest that MCH neuron activation-induced REM sleep is indistinguishable from physiologically occurring REM sleep in terms of EEG, EMG, and eye and whisker movement indicators.

Activation of MCH neurons during NREM sleep but not during wakefulness induces REM sleep

The results presented in Figure 2 clearly demonstrate that REM sleep can be induced by MCH neuron stimulation during NREM sleep. To determine whether activation of MCH neurons during wakefulness also induces a transition to REM sleep, MCH neurons were intermittently activated for 1 min (475 ± 17.5 nm, 25.8 mW/mm², 10 ms, 10 Hz) every 5 min from 20:00 (the onset time of the active dark period) to 16:46 (the second half of the inactive light period). *MCH-tTA*; *TetO Chr2* mice without light illumination were used as controls. Figure 3A shows a representative hypnogram from *MCH-tTA*; *TetO Chr2* mice under basal conditions (without light illumination). In comparison, the hypnogram of *MCH-tTA*; *TetO Chr2* mice with blue light illumination (Fig. 3B) shows frequent transitions to REM sleep in conjunction with light illumination. However, transitions to REM sleep were restricted to epochs during which mice were in NREM sleep. Activation of MCH neurons in NREM sleep immediately decreased EEG delta power and increased theta power, suggesting a transition to REM sleep (Fig. 3C,D). Mean REM sleep latency calculated from the initiation of blue light illumination was 12.4 ± 1.2 s ($n = 129$). REM sleep was often terminated when blue light illumination ended. In contrast, activation of MCH neurons did not induce a transition to REM sleep when mice were awake (Fig. 3E,F). EEG and EMG were indistinguishable from before MCH neuron activation.

Figure 3G–J presents the percentage of transitions during and after 1 min MCH neuron activation during the dark period (Fig.

3G,H) and in the light period (Fig. 3I,J). Activation of MCH neurons was initiated either when the mice were awake (Fig. 3G,I) or in NREM sleep (Fig. 3H,J). Activation of MCH neurons while the mice were awake during either the dark or light period did not affect vigilance states; wakefulness was maintained both during and after activation of MCH neurons (Fig. 3G,I). In contrast, activation of MCH neurons during NREM sleep greatly increased the probability of transition to REM sleep (Fig. 3H,J). REM sleep duration was ~ 1 min, probably due to the duration of light illumination. MCH neuron activation during NREM sleep in the light period was particularly effective in facilitating the transition from NREM to REM sleep (Fig. 3J): the proportion of mice transitioning from NREM sleep to REM sleep within 1 min of MCH neuron activation was $89.7 \pm 1.2\%$ ($n = 7$, $p < 0.001$, unpaired t test). To exclude the effect of blue light illumination alone on the transition to REM sleep, littermate monogenic *MCH-tTA* or *TetO Chr2* mice were subjected to blue light illumination. In these mice, the transition probability from NREM sleep to REM sleep after 1 min blue light MCH neuron illumination was only $10.4 \pm 1.6\%$ ($n = 8$). The EEG power spectral analysis by FFT showed that EEG power of 3–8 Hz was significantly increased during light illumination in the light period (8:00 to 16:46; Fig. 3K). The average EEG power density in the delta (1–5 Hz) and theta (6–10 Hz) wave bands was significantly increased as well during light illumination (Fig. 3L), indicating an increase in sleep intensity during activation of MCH neurons.

Acute inhibition of MCH neurons has no effect on sleep/wake states

Next, the effect of acute inhibition of MCH neurons on vigilance states was determined. To strongly inhibit MCH neurons over a long period using optogenetics, *TetO ArchT* mice (Tsunematsu et al., 2013) were bred with *MCH-tTA* mice to generate bigenic *MCH-tTA*; *TetO ArchT* mice (Fig. 4A).

The specific expression of ArchT in MCH neurons was confirmed by a double-label immunohistochemical study. An anti-GFP antiserum was used to detect ArchT-EGFP fusion proteins. A merged picture (GFP-immunoreactive and MCH-immunoreactive) revealed that ArchT-EGFP was exclusively expressed in MCH neurons (Fig. 4B,C). ArchT expression was restricted to MCH neurons; ectopic expression of ArchT outside of MCH neurons was not observed. Confocal microscopic observation revealed that ArchT was present in the soma and dendrites of MCH neurons. The number and morphology of MCH-immunoreactive neurons in *MCH-tTA*; *TetO ArchT* mice were indistinguishable from those of monogenic littermate mice (*MCH-tTA* or *TetO ArchT* mice; data not shown), suggesting that ArchT expression is not toxic to MCH neurons. ArchT was expressed in $97.3 \pm 2.0\%$ ($n = 6$, GFP-immunoreactive/MCH-immunoreactive $\times 100\%$) of MCH-immunoreactive neurons.

Slice patch-clamp recording from MCH neurons confirmed the function of ArchT in MCH neurons. Under whole-cell current-clamp mode, a rectangular depolarizing current (0.2 Hz, 100 ms) was injected through the recording electrode to generate artificial action potentials since most MCH neurons were silent. Green light illumination (549 ± 7.5 nm, 0.9 mW) for 1 min completely inhibited current injection-generated action potentials (Fig. 4D). Green light illumination completely inhibited generation of action potentials. The firing probability during green light illumination was $0.0 \pm 0.0\%$ ($n = 7$, $p < 0.001$; Fig. 4E). After termination of green light illumination, neuronal discharge rate recovered to $98.8 \pm 1.3\%$ of the preillumination firing rate ($n = 7$, $p = 0.24$, NS).

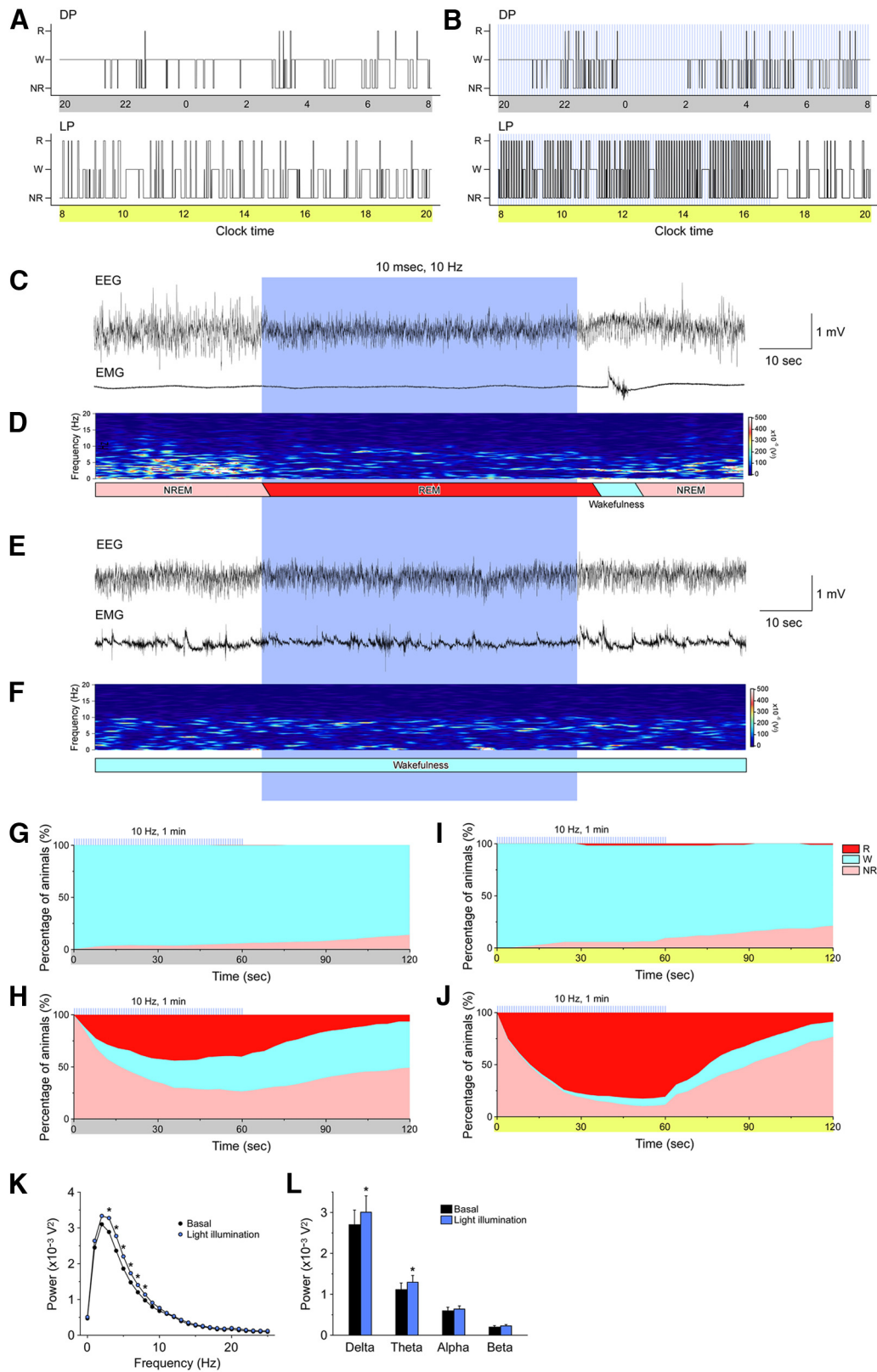


Figure 3. Acute activation of MCH neurons induces transitions from NREM sleep to REM sleep. **A, B**, Representative hypnograms of (**A**) no illumination and (**B**) blue light illumination for 1 min every 5 min (25.8 mW/mm^2 , 10 ms, 10 Hz) from 20:00 to 16:46. The upper panel is the dark period (20:00–8:00); lower panel is the light period (8:00–20:00); blue bars indicate 1 min periods of blue illumination. **C, E**, Representative traces for EEG (upper trace) and EMG (lower trace). Blue pulses were applied during NREM sleep (**C**) or wakefulness (**E**). **D, F**, EEG power spectra corresponding to **C** and **E**. **G–J**, Graphs show percentage of animals in each vigilance state during blue pulse illumination. **G, H**, Dark period. **I, J**, Light period. Blue light illumination occurred during a waking period in **G** and **I** and during NREM sleep in **H** and **J**. The blue lines above each graph indicate 1 min blue light illumination pulses. **K**, Power spectral analysis of EEG in the light period (8:00–16:46) without illumination (black) and with illumination (blue). **L**, The average EEG power densities in the delta, theta, alpha, and beta wave bands in the light period (8:00–16:46). W, Wakefulness; R, REM sleep; NR, NREM sleep; LP, light period; DP, dark period. Values are represented as means \pm SEM. * $p < 0.05$ versus no illumination.

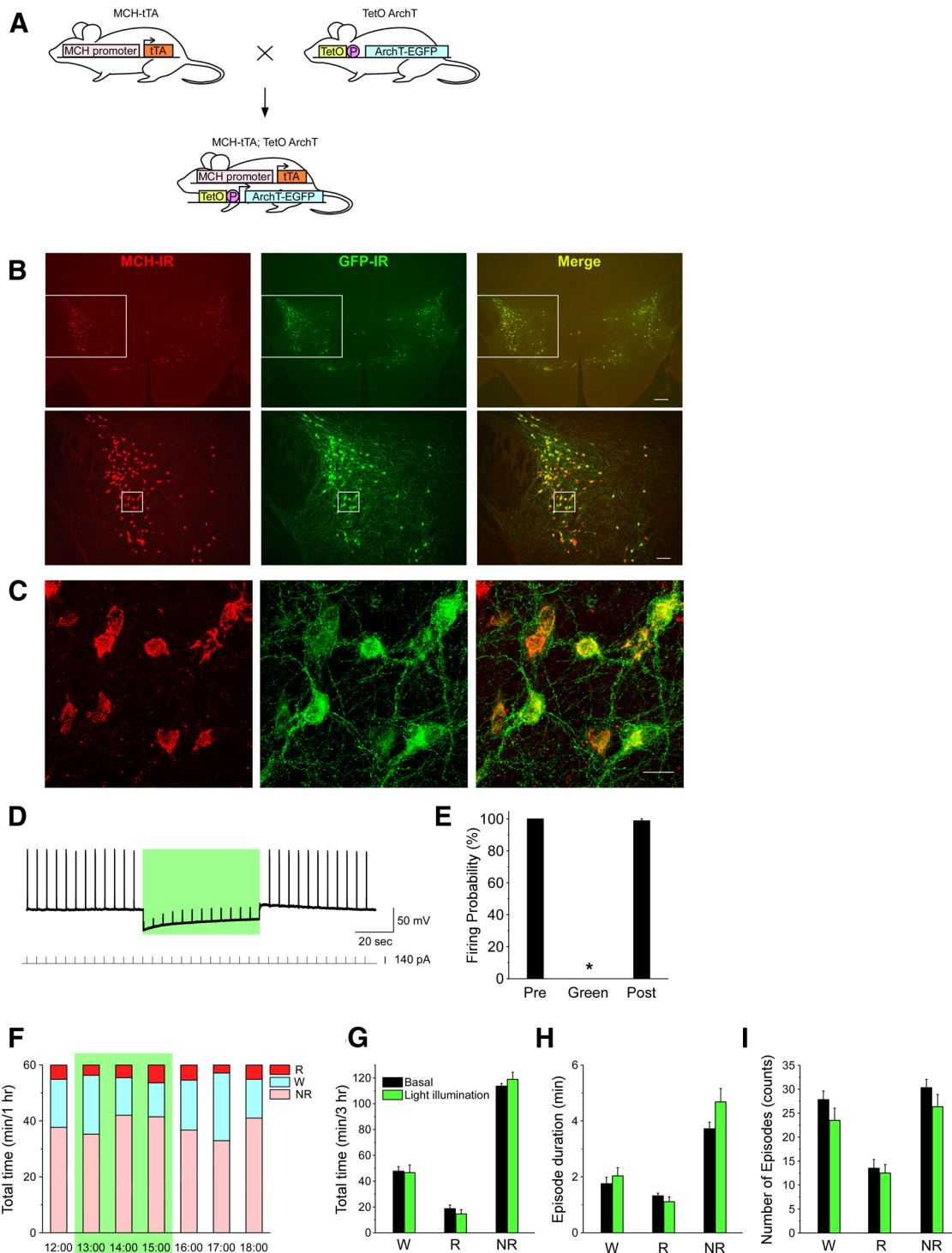


Figure 4. *In vivo* optogenetic inhibition of MCH neurons has no effect on vigilance states. **A**, Schematic shows generation of bigenic *MCH-tTA*; *TetO ArchT* mice. P, Minimal promoter. **B, C**, Immunohistochemical analyses revealed that ArchT is specifically expressed in MCH neurons in the *MCH-tTA*; *TetO ArchT* bigenic mouse brain. **B**, MCH-immunoreactive (MCH-IR) neurons (left, red; Alexa594) and ArchT::GFP-immunoreactive (GFP-IR) neurons (middle, green; Alexa488) located in the LHA and the zona incerta. Right, Merged image shows specific expression of ArchT in MCH neurons. Bottom row are higher magnifications of the regions enclosed by the squares in the top row. Scale bars: (in top, right) top row, 200 μ m; (in bottom, right) bottom row, 100 μ m. **C**, Confocal microscopic image of the region indicated by the squares in **B**, bottom row. Scale bar, 20 μ m. **D, E**, Slice patch-clamp recordings from MCH neurons. **D**, Upper trace, Current-clamp recording from MCH neuron. Lower trace, Injected current through the recording electrode (0.2 Hz, 140 pA, 100 ms) to generate action potentials. Green light illumination (549 \pm 7.5 nm, 0.9 mW, 1 min) inhibited firing elicited by current injection. **E**, Bar graph indicating the firing probability calculated from the data in **D** ($n = 7$). Firing probability is normalized to the 1 min before light illumination (Pre). Green light was applied for the duration indicated by the green bars. Values represent means \pm SEM. * $p < 0.05$ versus Pre. **F**, Green light illumination (542 \pm 13.5 nm, 33.4 mW/mm², 3 h from 13:00 to 16:00) in the light period. **G–I**, Bar graphs indicating the total time in each vigilance state (**G**), episode duration (**H**), and number of episodes (**I**) during green light illumination. W, Wakefulness; R, REM sleep; NR, NREM sleep. Values are represented as means \pm SEM. * $p < 0.05$ versus basal (no illumination).

During the light (inactive) period, green light (542 ± 13.5 nm, 33.4 mW/mm²) was illuminated through fiber optics onto the hypothalamus of *MCH-tTA*; *TetO ArchT* mice for 3 h from 13:00 to 16:00 using the same methods used for activation of MCH neurons. *MCH-tTA*; *TetO ArchT* mice without illumination were used as controls. Optogenetic inhibition of MCH neurons had no significant effect on time spent in any vigilance state (Fig. 4*F,G*). In the control condition, the time spent in wakefulness, REM sleep, and NREM sleep over the 3 h period was 47.7 ± 3.5 min ($n = 6$), 18.6 ± 2.9 min ($n = 6$), and 113.6 ± 2.1 min ($n = 6$), respectively. The time spent in wakefulness, REM sleep, and NREM sleep during green light illumination was 46.6 ± 5.9 min ($n = 6$, $p = 0.88$, NS, paired *t* test), 14.5 ± 3.3 min ($n = 6$, $p = 0.054$, NS, paired *t* test), and 118.9 ± 5.5 min ($n = 6$, $p = 0.41$, NS, paired *t* test), respectively. Inhibition of MCH neurons also had no effect on episode duration (Fig. 4*H*) or the number of episodes (Fig. 4*I*). As indicated in Table 1, ArchT expression in MCH neurons had no effect on basal sleep/wakefulness patterns (total time and episode duration of each state). These results suggest that silencing of MCH neurons has little effect on the initiation and maintenance of REM sleep.

Ablation of MCH neurons decreases time spent in NREM sleep

Inhibition of MCH neurons using optogenetics had little effect on sleep/wakefulness. However, due to the technical limitations of optogenetic inhibition, particularly for long periods of inhibition, we wanted to induce a chronic loss of function to further evaluate the role of MCH neurons in sleep/wakefulness regulation. Thus, we used the tet-off system mice to specifically ablate MCH neurons. *MCH-tTA* mice were bred with *TetO DTA* mice to generate *MCH-tTA*; *TetO DTA* bigenic mice (Fig. 5*A*). In these mice, DTA expression is restricted to MCH neurons and DTA induces cell death by inhibiting protein synthesis. The timing of DTA expression is controlled by the presence or absence of Dox in the chow. In the presence of DTA, tTA loses its ability to bind the TetO sequence (Fig. 5*B*). However, in the absence of Dox, tTA induces DTA expression. Here, *MCH-tTA*; *TetO DTA* mice were fed with chow containing Dox (100 mg/kg) until 10 weeks of age. Then, Dox(+) chow was replaced with Dox(-) chow, the timing of which was defined as Dox(-) 0 week (Fig. 5*C*).

MCH neuron-specific and temporally controlled ablation was confirmed by immunohistochemical studies using an anti-MCH antibody. The number of MCH-immunoreactive neurons was counted in C57BL/6J wild-type mice, in *TetO DTA* monogenic mice at 10 weeks of age, and in *MCH-tTA*; *TetO DTA* bigenic mice at DOX(-) 0, 1, 2, 3, and 4 weeks. At Dox(-) 0 week, the number of MCH neurons was 1028 ± 33 ($n = 3$). This was comparable to the number of MCH neurons in the C57BL/6J wild-type mice (1076 ± 61 , $n = 4$, $p = 1$, Kruskal–Wallis) and in the *TetO DTA* monogenic mice (1031 ± 65 , $n = 3$, $p = 1$, Kruskal–Wallis). Meanwhile, the number of MCH neurons dramatically decreased after Dox removal (Fig. 5*D,E*). The number of MCH neurons at Dox(-) 1, 2, 3, and 4 weeks was 600 ± 52 (58.3%, $n = 3$, $p < 0.001$, Kruskal–Wallis), 112 ± 18 (10.9%, $n = 4$, $p < 0.001$, Kruskal–Wallis), 55 ± 4 (5.4%, $n = 3$, $p < 0.001$, Kruskal–Wallis), and 28 ± 9 (2.7%, $n = 4$, $p < 0.001$, Kruskal–Wallis), respectively. To confirm whether DTA-induced cell death induction was restricted to MCH neurons, orexin/hypocretin neurons were stained and counted (Fig. 5*F,G*). Orexin neurons are distributed in the lateral hypothalamic area but do not colocalize with MCH (Broberger et al., 1998; Elias et al., 1998; Peyron et al., 1998; Bayer et al., 2002). The number of orexin

neurons was comparable to that of C57BL/6J wild-type mice and that of *TetO DTA* monogenic mice (Fig. 5*G*). In addition, the number of orexin neurons was unaffected by Dox removal for 4 weeks in *MCH-tTA*; *TetO DTA* mice (Fig. 5*G*). No signs of unhealthiness, shape, or size of orexin neurons were found even though orexin neurons are located near MCH neurons. These results confirm that the tet-off system functioned correctly in the transgenic mice and that MCH neurons were specifically ablated after removing Dox from the chow.

Next, the effect of ablation of MCH neurons on sleep/wakefulness regulation was analyzed. EEG and EMG electrodes were implanted in *MCH-tTA*; *TetO DTA* mice at 8 weeks of age and EEG and EMG recordings were started after 2 weeks recovery at 10 weeks of age.

MCH-tTA; *TetO DTA* mice, fed with Dox(+) chow [the Dox(+) group], were used as controls. The total time in wakefulness, REM sleep, and NREM sleep was unaffected in the Dox(+) group during recordings [from Dox(+) 0 week to Dox(+) 4 weeks]. In contrast, the time spent in wakefulness was significantly increased and the time in NREM sleep decreased in the Dox(-) group (Fig. 5*H*). At Dox(-) 4 weeks, total time in wakefulness in the Dox(+) group and the Dox(-) group was 697.1 ± 29.6 min ($n = 5$) and 793.1 ± 13.5 min ($n = 8$, $p < 0.001$, unpaired *t* test), respectively. At Dox(-) 4 weeks, the total time in NREM sleep in the Dox(+) group and the Dox(-) group was 664.2 ± 30.5 min ($n = 5$) and 572.7 ± 14.8 min ($n = 8$, $p = 0.01$, unpaired *t* test), respectively. Although increases in wakefulness were observed during both the light and dark periods, the increase was more pronounced in the light period. Interestingly, however, the time spent in REM sleep was unaffected in either the light or dark period (Fig. 5*He,Hf*).

The mean episode duration of NREM sleep across the 24 h period and during the dark period significantly decreased from Dox(-) 2 weeks (Fig. 5*Ig,Ii*). The mean episode durations of wakefulness and REM sleep were not significantly affected (Fig. 5*Ia–Ij*). These results provide evidence that MCH neurons are involved in the regulation of NREM sleep in addition to their role in REM sleep demonstrated above.

To confirm that the ablation of MCH neurons did not affect normal cortical activity, cortical EEG power spectral analyses were compared between Dox(-) 0 week and Dox(-) 4 weeks in *MCH-tTA*; *TetO DTA* mice (Fig. 6). The power spectra in both REM and NREM sleep of Dox(-) 4 weeks mice were indistinguishable from those of Dox(-) 0 week mice, suggesting that ablation of MCH neurons had no effect on basal cortical activity.

Discussion

In the present study, newly developed transgenic mouse strains enabled manipulation of the activity and fate of MCH neurons *in vivo*. The results obtained support a role for MCH neurons in the regulation of both REM and NREM sleep.

Acute and reversible manipulation of MCH neuronal activity using optogenetics

A role for MCH neurons in sleep/wake regulation has previously been suggested based on studies using gene knock-out mice and pharmacological studies. *Prepro-MCH* gene knock-out mice exhibited longer wake bouts during the dark period but REM sleep was unchanged (Willie et al., 2008). *MCHR1* gene knock-out mice increased wakefulness and decreased NREM sleep (Ahnaou et al., 2011). *MCHR1* antagonists decreased deep NREM and REM sleep quantities primarily by reducing the mean episode duration (Ahnaou et al., 2008). MCH neurons are silent during

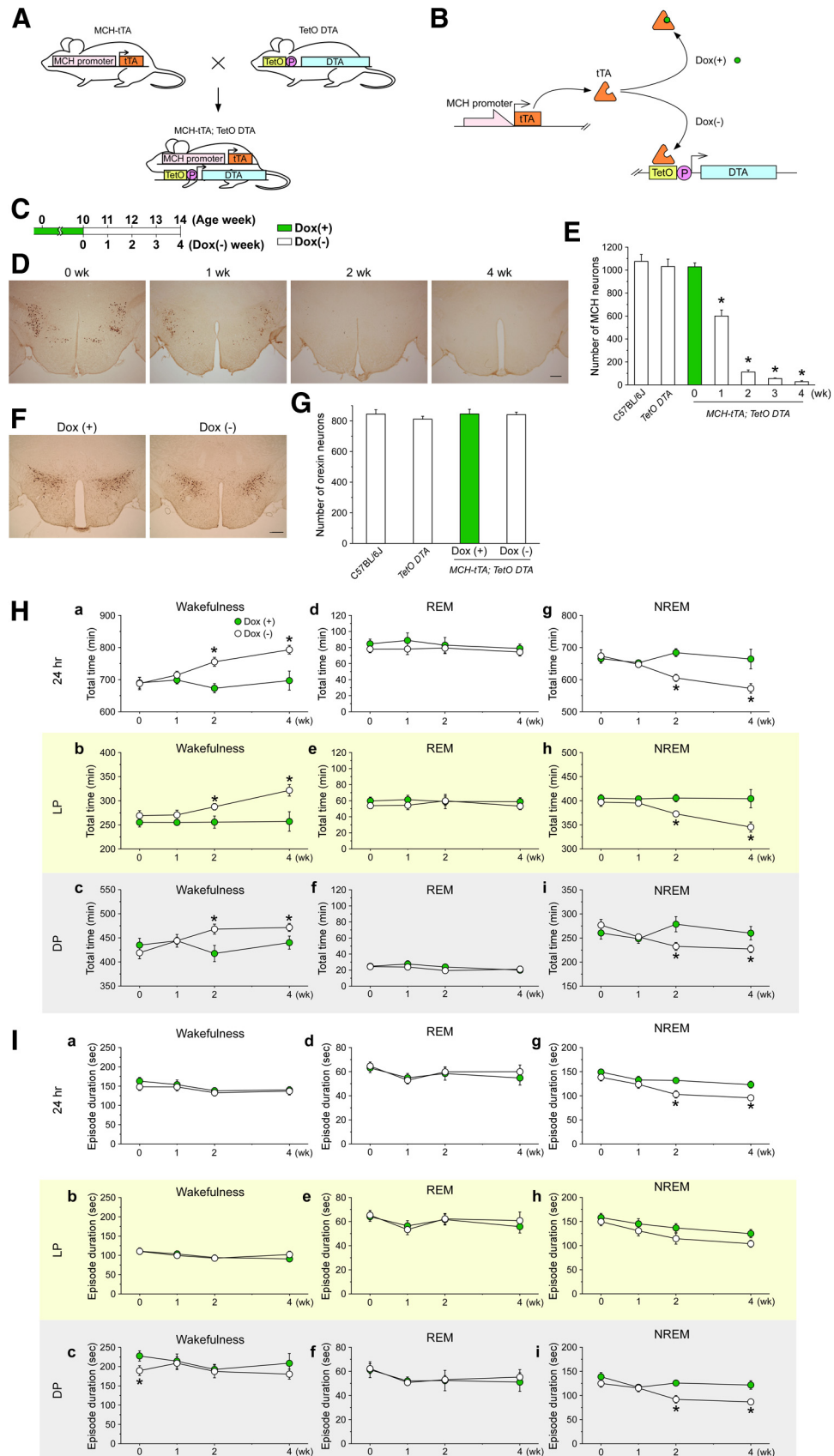


Figure 5. Ablation of MCH neurons increases wakefulness and decreases NREM sleep in *MCH-tTA; TetO DTA* bigenic mice. **A**, Schematic showing generation of bigenic *MCH-tTA; TetO DTA* mice. **B**, Minimal promoter. **B**, Schematic illustration of the tetracycline-controlled gene expression system and tTA-induced DTA expression in MCH neurons. In the presence of Dox, Dox binds tTA protein and DTA expression is repressed. **C**, Schematic shows the experimental protocol. In the presence of Dox, DTA expression is suppressed and MCH neurons are intact. Green bar, Chow contained Dox [Dox(+)]. Open bar, Chow without Dox [Dox(-)]. **D**, Immunohistochemistry indicates MCH neuron-specific ablation in Dox(-); 0, 1, 2, and 4 weeks (wk) (Figure legend continues.)

waking and discharge during sleep and are maximally active during REM sleep (Hasani et al., 2009). Despite these reports, however, the specific role of MCH neurons in sleep/wakefulness regulation remains uncertain.

To help clarify the role of MCH neurons in sleep/wake control, we determined the effects of acute optogenetic activation of MCH neurons. To activate these cells, we used a ChR2 (E123T/T159C) double mutant that is suitable for reliable and fast optical stimulation (Berndt et al., 2011). We found that acute activation of MCH neurons significantly increased the time spent in REM sleep in association with decreased NREM sleep. These results were in good agreement with those of a previous study in which ChR2 was expressed in MCH neurons using a Cre-dependent viral vector in *prepro-MCH-cre* transgenic mice (Jego et al., 2013). Interestingly, NREM rebound was observed after stimulation of MCH neurons. This might be caused simply by NREM suppression during activation of MCH neurons. Alternatively, this might suggest that homeostatic pressure for NREM sleep accumulates not only in wakefulness, but also in REM sleep.

On the other hand, optogenetic activation of MCH neurons has also been reported to increase both NREM and REM sleep (Konadhode et al., 2013). This discrepancy may be due to differential ChR2 expression among MCH neurons. The present study and that of Jego et al. (2013) report that >80% of MCH neurons expressed ChR2, whereas Konadhode et al. targeted MCH neurons that were dorsal to the fornix (53%) with relatively little expression (~20%) in the lateral and ventral hypothalamic populations (Konadhode et al., 2013). Activation of a subset of MCH neurons might have distinct effects on NREM sleep. Jego et al. (2013) expressed ChR2, eNpHR3.0, or ArchT in MCH neurons by viral injections into Cre transgenic mice. However, the expression rate among individual mice could differ since this technique

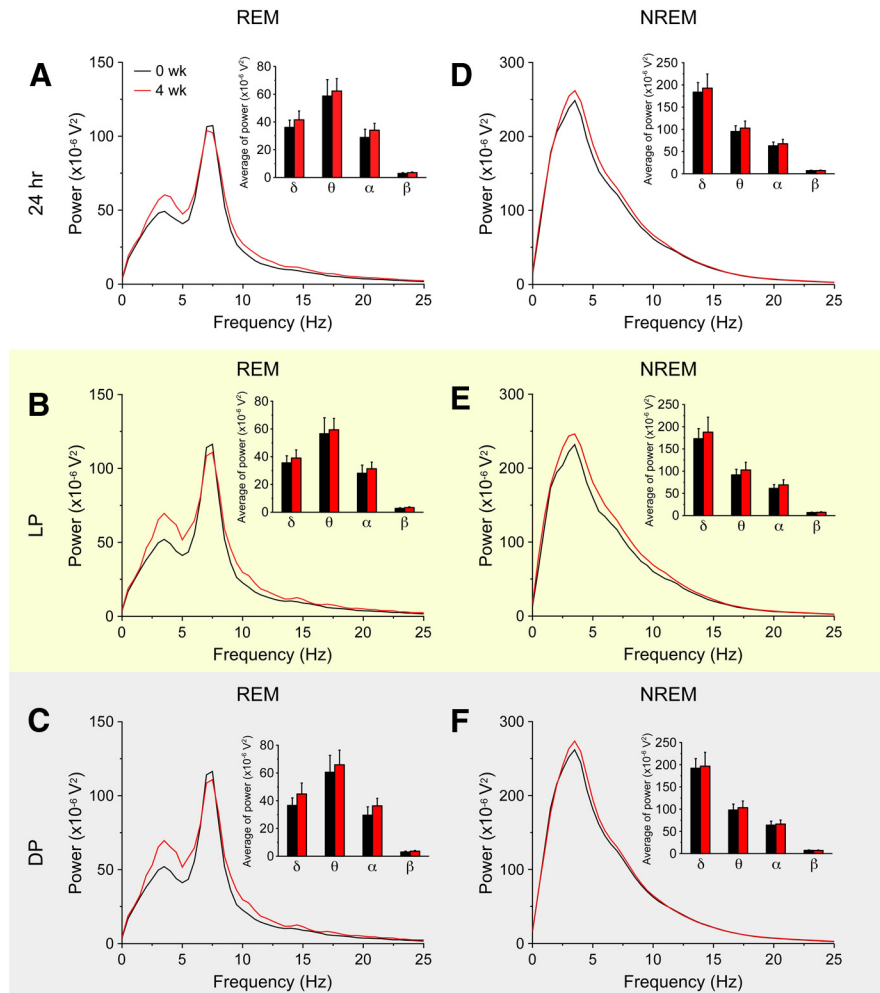


Figure 6. Ablation of MCH neurons had no effect on EEG power spectra during REM sleep and NREM sleep. **A–F**, Power spectral analysis of EEG recorded from *MCH-tTA; Teto DTA* bigenic mice. Black lines indicate the presence of Dox (MCH neurons are intact); Red lines show Dox removal for 4 weeks (MCH neurons are ablated). Inset summarizing the average EEG power density in the delta (1–5 Hz), theta (6–10 Hz), alpha (10–13 Hz), and beta (13–25 Hz) bands during REM sleep (**A–C**) and NREM sleep (**D–F**). Values are represented as means \pm SEM. * $p < 0.05$ versus 0 week.

is dependent upon interanimal replicability of each virus injection. In contrast, the bigenic mouse strains used in the present study do not require viral injections and we have found interanimal expression to be reliable and reproducible.

We further demonstrated that acute activation of MCH neurons during NREM sleep reliably triggered transitions to REM sleep. MCH neuron activation-induced REM sleep following NREM sleep suggests a similarity to physiological REM sleep control. In contrast, optogenetic silencing of MCH neurons had no effect on either total REM time or the duration of REM sleep episodes. These results demonstrate that MCH neuronal activity is sufficient to induce REM sleep, but is not necessary for REM sleep occurrence. This conclusion is consistent with the observation that REM sleep occurs even in pontine cats in which all structures rostral to the brainstem have been removed (Jouvet, 1962), implicating neurons in the brainstem as responsible for the generation of REM sleep.

MCH neurons may cooperatively interact with other brain regions to regulate REM sleep. Recent studies using cFos immunostaining with retrograde tracing have revealed an interaction between MCH neurons and pontine sublaterodorsal tegmental nucleus (SLD) neurons (Clément et al., 2012; Luppi et al., 2013a).

←

(Figure legend continued.) indicates time after Dox(–). Scale bar, 300 μ m. **E**, Bar graph shows the number of MCH-immunoreactive neurons ($n = 3–4$). **F**, Orexin neurons were stained with anti-orexin antibody to produce a golden-brown color (scale bar, 300 μ m). **G**, Bar graph summarizing the number of orexin-immunoreactive cell bodies in the LHA. **H**, Line graphs indicate the total time of each vigilance state. **Ha, Hd, Hg**, Total time in wakefulness, REM sleep, and NREM sleep, respectively. **Hb, He, Hh**, Total time for 24 h during the light period in wakefulness, REM sleep, and NREM sleep. **Hc, Hf, Hi**, Total time during the dark period in wakefulness, REM sleep, and NREM sleep. **I**, Line graphs indicate the mean duration of each vigilance state. **Ia, Id, Ig**, Mean duration over 24 h in wakefulness, REM sleep, and NREM sleep, respectively. **Ib, Ie, Ih**, Mean duration during the light period in wakefulness, REM sleep, and NREM sleep, respectively. **Ic, If, Ii**, Mean duration during the dark period in wakefulness, REM sleep, and NREM sleep. Green circles, Dox(+) condition (MCH neurons are intact); open circles Dox(–) (MCH neurons are ablated). Values are represented as means \pm SEM. * $p < 0.05$ versus Dox(+). LP, Light period; DP, dark period.

Other studies indicate that REM sleep is generated by glutamatergic neurons in the SLD that are specifically active during REM sleep (Sakai and Koyama, 1996; Boissard et al., 2002; Lu et al., 2006; Clément et al., 2011). These studies also suggest that MCH neurons inhibit GABAergic REM-off neurons located in the ventrolateral periaqueductal gray region during REM sleep and that these GABAergic REM-off neurons gate the activation of the REM-on glutamatergic neurons in the SLD. MCH neurons might function to open this gate to initiate REM sleep during NREM sleep. This hypothesis is supported by our results following intermittent activation of MCH neurons. Although activation of MCH neurons during NREM sleep induced REM, activation during wakefulness had no effect on either REM or NREM sleep.

Direct transitions from wakefulness to REM sleep do not occur under normal conditions, but it occurs in narcoleptics in whom orexin signaling pathways are interrupted. This observation indicates that, in contrast to MCH, orexin inhibits the initiation of REM sleep. Indeed, local administration of orexin into the locus ceruleus or the laterodorsal tegmental nucleus dramatically suppresses REM sleep without affecting NREM sleep quantity (Bourgin et al., 2000; Xi et al., 2001). Orexin neurons and MCH neurons are coextensive in the LHA, which leads us to imagine a functional interaction between these neurons in the regulation of sleep/wakefulness. Our observation that MCH neuron activation during NREM sleep induces REM sleep with higher probability in the light period compared with the dark period might reflect this interaction. Orexin concentrations in the CSF are higher during the dark period compared with the light period (Yoshida et al., 2001), which might interfere with the transition from NREM to REM sleep when MCH neurons are activated.

Chronic and irreversible ablation of MCH neurons

To further elucidate the role of MCH neurons in sleep/wakefulness, MCH neuron-specific ablation was performed. Near-complete MCH neuron ablation increased the total time in wakefulness with decreased NREM sleep. However, the total time in REM sleep was not affected. These observations indicate that MCH neurons also play a crucial role in the regulation of NREM sleep. The result from this chronic disruption of MCH neurotransmission conflicted with our results demonstrating that acute activation of MCH neurons increases total time in REM sleep. How could such different results between chronic ablation and acute activation of MCH neurons arise? One possible explanation is suggested by a slice patch-clamp study that revealed that MCH exhibited an inhibitory effect on orexin neurons (Rao et al., 2008). The orexin and MCH systems often display opposing effects on physiological functions. For example, MCH knock-out mice were hyperactive during fasting (Willie et al., 2008), whereas orexin neuron-ablated mice were hypoactive but did not increase locomotor activity during fasting (Hara et al., 2001; Yamanaka et al., 2003). Orexin neurons are thought to be active during wakefulness and silent during REM sleep (Lee et al., 2005; Mileykovskiy et al., 2005; Hassani et al., 2009), whereas MCH neurons are active in REM sleep and inactive during wakefulness (Hassani et al., 2009). Putative somasomatic, axosomatic, and axodendritic contacts between orexin and MCH neurons have been observed (Bayer et al., 2002; Guan et al., 2002). Additionally, MCHR1 knock-out mice showed an increase in glutamatergic transmission onto orexin neurons. These facts support the concept that MCH neurons have a role in the inhibition of orexin neuronal activity. Therefore, ablation of MCH neurons might result in a chronic activation of orexin neurons, which might cause an in-

crease in the total wake time and decrease in NREM sleep in MCH neuron-ablated mice. However, acute inhibition of MCH neurons affected neither the time in wakefulness nor NREM sleep duration. Optogenetic inhibition for 3 h might not be enough to affect orexin neurons. Alternatively, after ablation of MCH neurons, neural network reorganization might occur, resulting in chronic activation of orexin neurons.

MCH neurons also project to forebrain areas, such as the hippocampus and cortex (Bittencourt et al., 1992; Saito et al., 1999; Hervieu et al., 2000). High levels of MCHR1 mRNA and MCHR1 immunoreactivity have been found in these areas (Bittencourt et al., 1992; Saito et al., 1999; Hervieu et al., 2000). Therefore, in addition to its role in sleep/wakefulness regulation, the release of MCH onto cortical and hippocampal cells during REM sleep might modulate long-term plasticity and contribute to the processing of learning and memory. Similar to the result of Konadhode et al., the increase of delta and theta power during activation of MCH neurons might suggest the role of MCH on the cortex (Konadhode et al., 2013). Additional studies are also required to determine the role of MCH release during REM sleep in these brain areas.

Together, the present results show that MCH neurons in the hypothalamus play a significant role in the regulation of both NREM and REM sleep. Further experiments are necessary to identify the specific neural pathways that underlie the differential roles that MCH neurons play in the regulation of NREM versus REM sleep. The creation of *MCH-tTA*; *TetO DTA* mice, which enable partial lesions of the MCH neuron population to be produced by varying the duration of Dox removal from the diet, may be a useful tool in this regard.

Notes

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