

# Octopamine Regulates Sleep in *Drosophila* through Protein Kinase A-Dependent Mechanisms

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Sleep is a fundamental process, but its regulation and function are still not well understood. The *Drosophila* model for sleep provides a powerful system to address the genetic and molecular mechanisms underlying sleep and wakefulness. Here we show that a *Drosophila* biogenic amine, octopamine, is a potent wake-promoting signal. Mutations in the octopamine biosynthesis pathway produced a phenotype of increased sleep, which was restored to wild-type levels by pharmacological treatment with octopamine. Moreover, electrical silencing of octopamine-producing cells decreased wakefulness, whereas excitation of these neurons promoted wakefulness. Because protein kinase A (PKA) is a putative target of octopamine signaling and is also implicated in *Drosophila* sleep, we investigated its role in the effects of octopamine on sleep. We found that decreased PKA activity in neurons rendered flies insensitive to the wake-promoting effects of octopamine. However, this effect of PKA was not exerted in the mushroom bodies, a site previously associated with PKA action on sleep. These studies identify a novel pathway that regulates sleep in *Drosophila*.

**Key words:** *Drosophila*; octopamine; sleep; locomotion; norepinephrine; biogenic amine; arousal

## Introduction

Sleep is a core process that spans genetically diverse eukaryotes from mammals to arthropods (Tobler, 2005). Disrupting sleep in any of these organisms is detrimental to their performance, memory, and health (Rechtschaffen, 1998). Extreme loss of sleep can even lead to death (Shaw et al., 2002). Thus, it must serve a very important function. Its conservation over evolution supports this claim for another reason: the need to sleep must outweigh selection pressure to eliminate it as a risk to predation. However, the function of sleep is unknown, and the molecular regulation underlying it is poorly understood.

Sleep can be monitored through electroencephalograms and electromyograms, but, when such electrophysiological recordings are technically difficult, as in the case of *Drosophila*, it is monitored through analysis of behavior. *Drosophila* show a sleep state characterized by changes in position, increased arousal threshold, and periods of inactivity that can last several hours (Hendricks et al., 2000; Shaw et al., 2000). Although little is known about the regulation of this sleep state, effects of some neurotransmitters have been described. Thus, GABA and serotonin promote sleep, the latter by acting through the 5-HT<sub>1A</sub> receptor expressed in the mushroom body (MB) (Yuan et al., 2006).

The only arousal-promoting signal identified in *Drosophila* is dopamine (Andretic et al., 2005; Kume et al., 2005).

In mammals, dopamine and norepinephrine are associated with states of arousal (Aston-Jones and Bloom, 1981). However, results regarding the effects of norepinephrine on total sleep and wake amounts have been mixed (Hunsley and Palmiter, 2003; Ouyang et al., 2004), in part because of differences in experimental protocols and also perhaps attributable to the effects of norepinephrine manipulation on dopamine levels (Schank et al., 2006). The insect equivalent of norepinephrine, octopamine, is synthesized and regulated through pathways that are distinct from those that produce dopamine. However, although octopamine is known to play a role in memory formation, larval locomotion, wing beating, ovulation, and aggression (Roeder, 2005), it has not been examined for effects on sleep. Here we demonstrate a novel role for octopamine in the regulation of sleep and wake in *Drosophila*. Feeding octopamine to flies leads to a protein kinase A (PKA)-dependent decrease in total sleep, whereas removal of octopamine from the food is followed by a sleep rebound. In addition, flies mutant for octopamine show an increase in total sleep, which can be restored to control levels with the administration of octopamine. We show that electrical excitation of octopamine-producing cells decreases total sleep, whereas electrical silencing of these cells increases sleep. Other parameters of sleep such as sleep latency and arousal threshold are also altered. Last, we demonstrate an activity-promoting role for the octopamine precursor tyramine, which is independent of the effects of octopamine.

## Materials and Methods

**Fly strains used.** Wild-type fly strains included *w;RC1;RC1* (isogenized chromosome 1 from the *w*<sup>1118</sup> stock; isogenized chromosomes 2 and 3 from the *RC1* strain) and *w*<sup>1118</sup>, *Canton S*, *Iso31* (isogenized *w*<sup>1118</sup> strain).

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Octopamine production mutants included *Tdc2*<sup>RO54</sup> (tyrosine decarboxylase 2 mutant) and *TβH*<sup>nm18</sup>/*FM7* (tyramine β hydroxylase mutant). Gal4 lines included *Tdc2-Gal4* (neuronal *Tdc2* expression pattern), *Tdc1-Gal4* (non-neuronal *Tdc1* expression pattern), *ElavGeneSwitch* (pan-neuronal expression during adulthood), and *MBSwitch* (mushroom body-specific expression during adulthood). Upstream activating sequence (*UAS*) lines included *UAS-B16B* (*NaChBac*) (bacterial Na<sup>+</sup> channel), *UAS-Kir2.1* (inward rectifying K<sup>+</sup> channel), *UAS-BDK33* (*PKAr*) (*Drosophila* inhibitory subunit of PKA with mutated cAMP binding site), *UAS-Tdc2*, and *UAS-Tdc1*. The following lines were ordered from the Bloomington Stock Center: *Tdc2-Gal4* (9313), *Tdc1-Gal4* (9312), *UAS-NaChBac* (9466), *UAS-Tdc2* (9315), *UAS-Tdc1* (9314), *UAS-GFPnls* (7032), and *Iso31* (5905). *UAS-Kir2.1* and *UAS-B16B* were a gift from Dr. B. White (National Institutes of Health, Bethesda, MD). *ElavGeneSwitch* (Osterwalder et al., 2001), *MBSwitch* (Mao et al., 2004), and *UAS-BDK33* (Rodan et al., 2002) were used previously in the laboratory (Hendricks et al., 2001; Joiner et al., 2006). The wild-type isogenic line *w;RC1;RC1* was a gift from W. Joiner (University of Pennsylvania, Philadelphia, PA). *Tdc2*<sup>RO54</sup> lines and *w*<sup>1118</sup> background line were a gift from Dr. G. Schupbach (Princeton University, Princeton, NJ) and Dr. J. Hirsh (University of Virginia, Charlottesville, VA). The *TβH*<sup>nm18</sup>/*FM7*, *TβH*<sup>me6</sup>/*FM7*, and *Canton S* background control were a gift from Dr. E. Kravitz (Harvard University, Boston, MA).

All *Gal4* and *UAS* lines were outcrossed seven times into the *w;RC1;RC1* or *Iso31* background. The *w;RC1;RC1* background was chosen for the *Tdc2-Gal4*-expressing *Kir2.1* because it shows lower levels of nighttime sleep (supplemental Table 1, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material) and thus affords the potential to avoid a ceiling effect.

**Sleep analysis.** Sleep analysis was performed as described previously (Joiner et al., 2006). All flies were kept on a 12 h light/dark (LD) cycle at 25°C schedule unless otherwise noted. Female and male flies 4–8 d old were placed in 65 × 5 mm tubes containing 5% sucrose and 2% agar and entrained for 24–36 h before the sleep recording. Baseline sleep was determined by monitoring activity for at least 3 d with no disruptions in an LD cycle. Locomotor activity was monitored using the DAMS/Trikinetics system as described previously (Joiner et al., 2006). Sleep was defined as a 5 min bout of inactivity as described previously (Shaw, 2003; Joiner et al., 2006). Latency to sleep was defined as the time in minutes from the moment lights were turned off to the first bout of sleep. Sleep consolidation scores were generated based on the amount of fragmentation seen in sleep, as measured by brief awakenings and the length of sleep bouts.

**Arousal threshold.** Arousal threshold was measured at three times of the night (2 h after lights off, 6 h after lights off, and 10 h after lights off). Increasing levels of mechanical stimulation were applied to determine the minimum stimulus for arousal. The levels were then labeled weak, medium, and strong, with weak being the lowest level of stimulation and strong being the maximum. Animals were scored based on their response to these three levels (Hendricks et al., 2000).

**Feeding octopamine and tyramine.** Wild-type (*w;RC1;RC1*, *Iso31*) female animals were loaded into monitors as described above and given 24–36 h to acclimate. One day of baseline data were collected, and then, at the lights-on transition, the flies were transferred either to tubes containing 5% sucrose/2% agar plus 10 mg/ml octopamine or onto sucrose/agar alone. Octopamine, administered at 10 mg/ml orally, was shown previously to be optimal for rescuing egg laying (Monastirioti et al., 1996). A dose–response curve of *Iso31* flies on octopamine is shown in supplemental Figure 3A (available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). The animals were left for 3 d on or off octopamine and removed at the lights-on transition. Rebound was determined and analyzed as described previously for sleep deprivation (Joiner et al., 2006). The males were examined in the same manner; however, 10 mg/ml octopamine was fatal for them over 3 d so the octopamine concentration was reduced to 5 mg/ml (data not shown). Other concentrations of octopamine were tested, but 10 mg/ml was found to be the optimal amount for 3 d for wild-type females (supplemental Fig. 3A, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). A blue food assay was performed as described (Edgecomb et al., 1994) to ensure that the animals

were eating the food (data not shown). Because of the lethality seen with 10 mg/ml octopamine, the *TβH*<sup>nm18</sup> females and controls were placed on 7.5 mg/ml octopamine.

Tyramine was also fed to wild-type flies. Similar to what was seen with octopamine, tyramine at higher concentrations produced increased lethality so wild-type male flies were fed 5 mg/ml tyramine in 5% sucrose/2% agar. Females were placed on 10 mg/ml tyramine in 5% sucrose/2% agar. We tested other amounts of tyramine and determined that 10 mg/ml was the optimal concentration (data not shown).

Mianserin was also used to address octopamine signaling. Based on the work of Maqueira et al. (2005), who used this compound to block cAMP increases attributable to octopamine *in vitro*, it was used at 0.2 mg/ml. All mianserin experiments were done with *Iso31* control flies. Following the same protocol outlined above for octopamine, we placed flies on mianserin alone, mianserin plus 10 mg/ml octopamine, control food, or control plus 10 mg/ml octopamine.

**Hydroxyurea analysis.** Ablation of mushroom bodies with hydroxyurea (HU) was performed as described previously (de Belle and Heisenberg, 1994). First-instar larvae of *Iso31* flies were collected and placed on either a yeast paste and water mixture or a yeast paste and 50 mg/ml HU mixture for 4 h at 25°C. They were then washed with water and placed in regular food vials until adulthood. At 4–8 d after eclosion, animals were loaded into monitor tubes as described above. They were given 3 d of acclimation and then transferred onto 7.5 mg/ml octopamine, because high lethality was observed at 10 mg/ml. After 3 d, the animals were then transferred back onto normal 5% sucrose agar tubes. Sleep analysis was then performed as described above. After the completion of the sleep analysis, fly heads were dissected in 4% paraformaldehyde and fixed for 30 min, mounted on slides, and analyzed for loss of the α and β lobes of mushroom bodies.

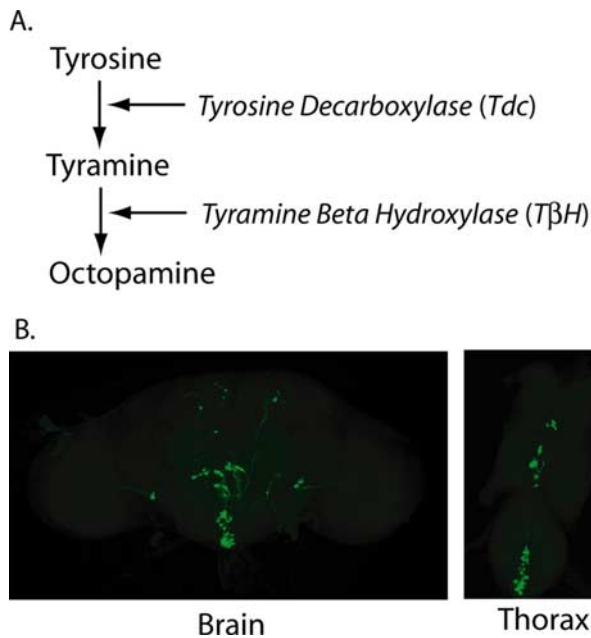
**PKA inhibition studies.** For PKA inhibition studies, we crossed the *ElavGeneSwitch* transgene into *UAS-BDK33* flies. Octopamine at 10 mg/ml was used as described above. Because the GeneSwitch construct can be turned on during adulthood using the drug RU486 (11β-(4-dimethylamino)phenyl-17β-hydroxy-17-(1-propynyl)estra-4,9-dien-3-one), we placed half the animals on 5% sucrose/2% agar tubes containing either 500 μM RU486 dissolved in ethanol or ethanol alone (1%) for 3 d. Half of each group was then transferred to octopamine-containing food with either 5% sucrose/2% agar plus 500 μM RU486 plus octopamine or 5% sucrose/2% agar plus ethanol (1%) plus octopamine. Both groups were also simultaneously fed 10 mg/ml octopamine for 3 d. At the lights-on transition at the end of this period, animals were transferred off of octopamine onto 5% sucrose/2% agar containing either 500 μM RU486 or ethanol (1%). Sleep analysis was performed as described above.

**Statistics.** To compare multiple groups, two-way ANOVA was used to determine significance for total sleep, nighttime sleep, daytime sleep, sleep bout number for both day and night, and latency to sleep. For non-Gaussian distributed data, we used the Kruskal–Wallis test; this included sleep bout length (daytime and nighttime), consolidation score, activity per waking minute, and peak activity. Statistical significance is denoted by asterisks: \**p* ≤ 0.01, \*\**p* ≤ .001, and \*\*\**p* ≤ .0001.

## Results

### Mutants with reduced octopamine have increased sleep

To address a role for octopamine and its precursor tyramine in *Drosophila* sleep, we analyzed two known genes that affect biosynthesis of these amines (Monastirioti et al., 1996; Cole et al., 2005; Certel et al., 2007). As shown in the flow diagram in Figure 1A, the *Tdc2* enzyme synthesizes tyramine (precursor to octopamine) from tyrosine; thus, its disruption results in low levels of octopamine and tyramine (Cole et al., 2005). In contrast, *TβH* synthesizes octopamine from tyramine, and therefore its loss reduces octopamine but increases tyramine levels 10-fold (Fig. 1A) (Monastirioti et al., 1996). Because synthesis of both amines requires *Tdc2*, the cellular distribution of *Tdc2* should reflect cells that potentially produce both tyramine and octopamine. We determined the expression pattern of *Tdc2* by using the well known

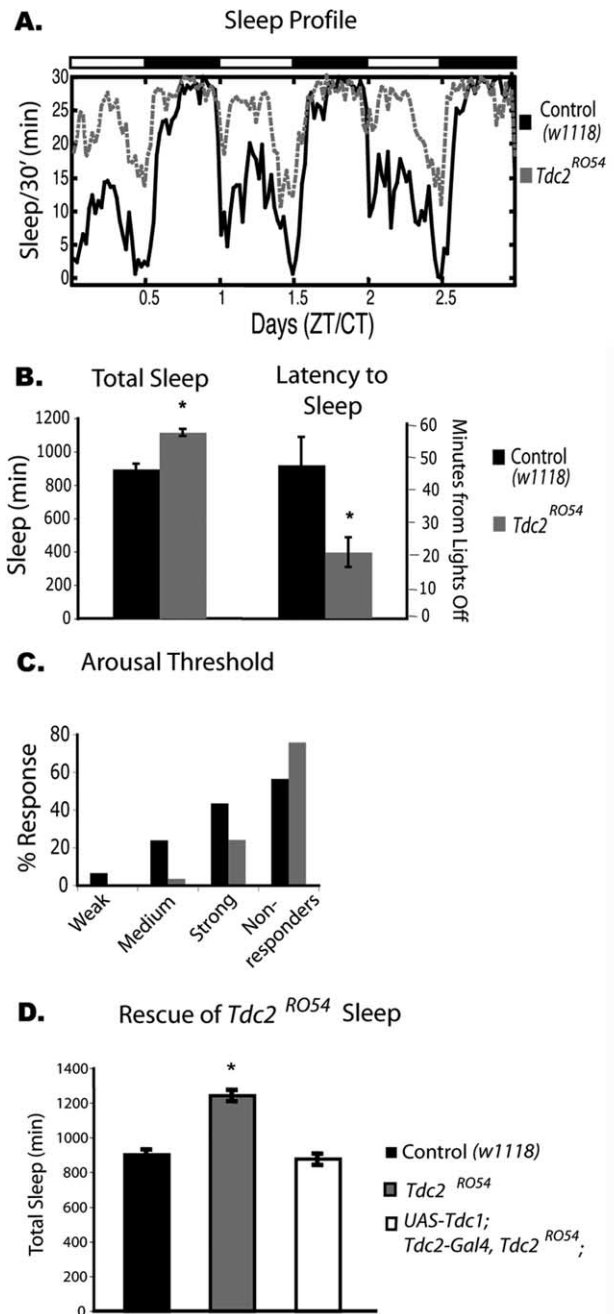


**Figure 1.** The octopamine biosynthesis pathway and its distribution in the fly brain. **A**, Octopamine and tyramine are derivatives of tyrosine. Tyrosine is converted to tyramine by *Tdc*. Tyramine is then converted to octopamine by *TβH*. **B**, Expression pattern of the *Tdc2-Gal4* line as visualized with a GFP reporter. Octopamine is produced in a subset of neurons in the brain and thoracic ganglion, respectively. Expression was characterized by crossing *Tdc2-Gal4* with nuclear *UAS-GFPnls*.

*UAS-Gal4* binary system to express green fluorescent protein (GFP) under the control of the *Tdc2* promoter (Fig. 1*B*). Expression of GFP was seen in discrete unilateral subsets of cells located along the ventral medial line of the brain, as well as in discrete bilateral clusters of cells in the lateral protocerebrum region and surrounding the esophagus cavity.

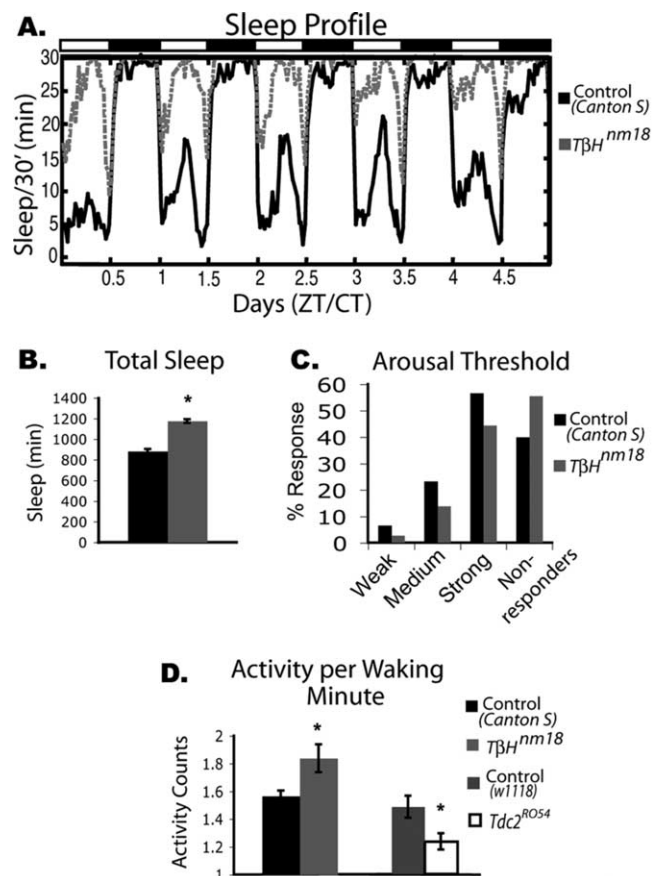
We then assayed sleep in flies mutant for each of the two genes in the octopamine biosynthesis pathway. Baseline sleep levels were examined in flies carrying a point mutation in the *Tdc* gene (*Tdc2<sup>RO54</sup>*) or a lesion, created by imprecise excision of a P transposable element, in the *TβH* gene (*TβH<sup>nm18</sup>*). Under baseline conditions, we found that male and female animals of both mutants displayed increased levels of sleep (Figs. 2*A, B*, 3*A, B*). All male data are shown in supplemental Table 1 (available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material), whereas female data are depicted in Figures 2–5. The increase in sleep in *Tdc2<sup>RO54</sup>* and *TβH<sup>nm18</sup>* mutants occurred primarily during the day (Figs. 2*A, B*, 3*A, B*), perhaps because flies are already sleeping maximally during the night. The increase in total sleep was accompanied by a decreased latency to sleep (the time from lights off until the animal's first bout of sleep) in the *Tdc2<sup>RO54</sup>* mutants (Fig. 2*B*), suggesting an increase in homeostatic drive to sleep. The *TβH<sup>nm18</sup>* mutation did not have a significant effect on latency, although there was a slight decrease in males (supplemental Table 1, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). The *Canton S* background strain that the *TβH<sup>nm18</sup>* mutants were crossed into displays rapid onset of sleep compared with other wild-type control lines used in this study (supplemental Table 1, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material), which may occlude any decrease in latency caused by the mutation.

Analysis of sleep architecture indicated that the increase in sleep in *Tdc2<sup>RO54</sup>* mutants was attributable to an increase in sleep bout number (supplemental Table 2, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material), whereas the increase in *TβH<sup>nm18</sup>*



**Figure 2.** Baseline sleep phenotype of *Tdc2<sup>RO54</sup>* mutants, which have decreased levels of octopamine and tyramine. **A**, Three days of baseline sleep recording in *Tdc2* females. The *Tdc2<sup>RO54</sup>* line (gray dashed line) shows significantly more sleep than its control *w<sup>1118</sup>* (black solid line). Dark bars and white bars on top indicate nighttime and daytime, respectively.  $n = 8$  for *Tdc2<sup>RO54</sup>* and *w<sup>1118</sup>*. CT, Circadian time; ZT, Zeitgeber time. **B**, Total sleep is significantly increased in *Tdc2<sup>RO54</sup>* mutant females (mean  $\pm$  SEM; *Tdc2<sup>RO54</sup>*,  $1115 \pm 19$ ,  $n = 23$ ; *w<sup>1118</sup>*,  $895 \pm 32$ ,  $n = 32$ ;  $p \leq 0.0001$ , two-way ANOVA). Latency to sleep is significantly lower in *Tdc2<sup>RO54</sup>* mutants (mean  $\pm$  SEM; *Tdc2<sup>RO54</sup>*,  $20.6 \pm 4.6$ ,  $n = 23$ ; *w<sup>1118</sup>*,  $47.5 \pm 8.7$ ,  $n = 32$ ;  $p \leq 0.001$ , two-way ANOVA). **C**, Arousal threshold during sleep in the *Tdc2<sup>RO54</sup>* mutants. The animals were given three levels of stimulation to determine whether they were arousable. All animals that responded to the first stimulation were also aroused on stronger stimulation. Compared with wild type, the *Tdc2* mutant line had a higher percentage of flies that did not respond to any of the three levels of stimulation. In addition, fewer *Tdc2<sup>RO54</sup>* flies responded to the weaker stimuli (*Tdc2<sup>RO54</sup>*,  $n = 23$ ; *w<sup>1118</sup>*,  $n = 32$ ). **D**, Rescue of the baseline sleep phenotype in *Tdc2<sup>RO54</sup>* with expression of the *Tdc1* gene. Total sleep was quantified, and *Tdc2<sup>RO54</sup>* mutant females were significantly different from both wild-type and rescued animals. There was no significant difference between rescued animals and controls (mean  $\pm$  SEM; *w<sup>1118</sup>*,  $895 \pm 33$ ,  $n = 32$ ; *Tdc2<sup>RO54</sup>*,  $1238 \pm 33$ ,  $n = 16$ ;  $p \leq 0.0001$ ; *UAS-Tdc1; Tdc2-Gal4, Tdc2<sup>RO54</sup>*,  $871 \pm 33$ ,  $n = 16$ ).





**Figure 3.** Baseline sleep phenotype of *TβH<sup>nm18</sup>* mutants, which have decreased levels of octopamine and increased levels of tyramine. **A**, Five days of baseline sleep recording in *TβH* females. The *TβH<sup>nm18</sup>* line (gray dashed line) shows significantly more sleep than its control, *Canton S* (black solid line). Dark bars and white bars on top indicate nighttime and daytime, respectively. Sixteen animals are shown for each group. CT, Circadian time; ZT, Zeitgeber time. **B**, Total sleep is significantly increased in the *TβH<sup>nm18</sup>* mutants (mean  $\pm$  SEM; *TβH<sup>nm18</sup>*,  $1176 \pm 17$ ,  $n = 46$ ; *Canton S*,  $884 \pm 20$ ,  $n = 52$ ;  $p \leq 0.0001$ , two-way ANOVA). **C**, Arousal threshold for the *TβH<sup>nm18</sup>* mutants. The animals were given three levels of stimulation to determine whether they were arousable. All animals that responded to the first stimulation were also aroused on stronger stimulation. Compared with wild type, the *TβH* mutant line had a higher percentage of flies that did not respond to any of the three levels of stimulation. In addition, fewer *Tdc2<sup>RO54</sup>* flies responded to the weaker stimuli (*TβH<sup>nm18</sup>*,  $n = 46$ ; *Canton S*,  $n = 52$ ). **D**, Activity per waking minute, which quantifies activity during the time the animal is awake, was significantly increased in the *TβH<sup>nm18</sup>* mutants (mean  $\pm$  SEM; *TβH<sup>nm18</sup>*,  $1.84 \pm 0.13$ ; *Canton S*,  $1.57 \pm 0.05$ ;  $p \leq 0.01$ , Kruskal–Wallis test) but significantly decreased in the *Tdc2<sup>RO54</sup>* mutants (mean  $\pm$  SEM; *Tdc2<sup>RO54</sup>*,  $1.2 \pm 0.09$ ; *w<sup>1118</sup>*,  $1.5 \pm 0.08$ ;  $p \leq 0.01$ , Kruskal–Wallis test).

mutants occurred from an increase in bout length (supplemental Table 2, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). Because both mutants affect octopamine similarly, this difference in sleep architecture is most likely attributable to differences in levels of tyramine. We also determined the arousal threshold in both mutants by measuring their response to a stimulus of increasing intensity. The arousal threshold during sleep was higher in both sets of mutants than in wild-type flies (Figs. 2C, 3C). This suggests that these animals are in a deeper state of sleep than their controls.

An increase in sleep could result from the animals being sick and unable to move. To address this possibility, we measured locomotor activity in *Tdc2<sup>RO54</sup>* and *TβH<sup>nm18</sup>*. We found that *Tdc2<sup>RO54</sup>* and *TβH<sup>nm18</sup>* had peak activity levels that were not significantly different from those of controls (data not shown).

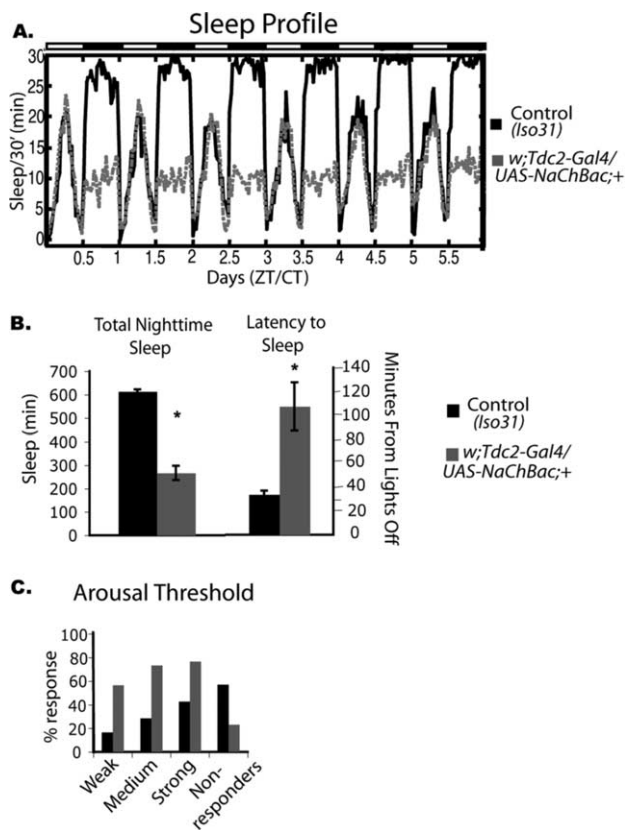
However, the *TβH<sup>nm18</sup>* flies showed significantly increased waking activity (Fig. 3D), as measured by total activity while awake; despite their increased total sleep time, the *TβH<sup>nm18</sup>* mutants showed a hyperactive phenotype when awake. In contrast, the average rate of movement was significantly decreased in *Tdc2<sup>RO54</sup>* flies compared with the wild-type controls (Fig. 3D). Because activity in the two mutants is affected in opposite directions, it is unlikely that the increased sleep phenotype of the mutants is secondary to effects on activity. More likely, the loss of octopamine (common to both mutants) underlies the decrease in sleep, whereas differences in tyramine levels account for effects on locomotor activity. In fact, our data are consistent with previously published data indicating that the increased (10-fold) levels of tyramine in the *TβH<sup>nm18</sup>* flies cause an increase in locomotor behavior when awake (Hardie et al., 2007).

It was shown previously that expression of *Tdc1* (the non-neuronal form of *Tdc*) in *Tdc2*-producing cells rescues the *Tdc2<sup>RO54</sup>* mutant locomotor phenotype (Hardie et al., 2007). We found the same to be true for sleep. When *Tdc1* was expressed in *Tdc2*-producing cells in a *Tdc2<sup>RO54</sup>* background, we found that we were able to rescue the baseline sleep phenotype (Fig. 2E), as well as sleep architecture and latency to sleep (data not shown).

#### Altering excitability of octopamine/tyramine-producing cells affects sleep

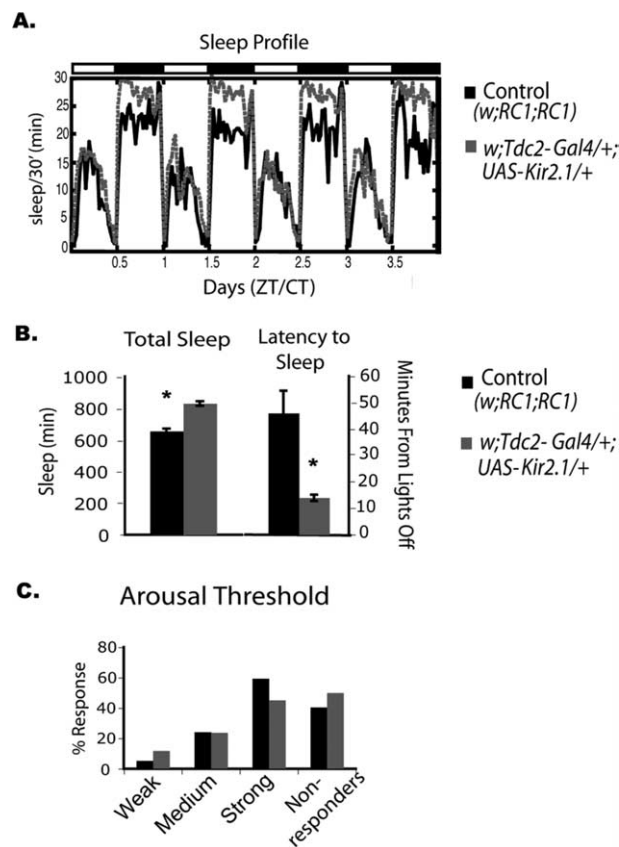
If octopamine and/or tyramine are released in the brain to regulate sleep, then blocking or increasing their release should also affect sleep. Thus, we sought to determine whether electrical manipulation of the cells producing tyrosine decarboxylase, which should affect the release of octopamine and tyramine, produces a change in sleep. The *Tdc2-Gal4* line mentioned above, which expresses Gal4 in cells producing octopamine and tyramine, was crossed to flies carrying transgenes for ion channels under the control of a *UAS* element recognized by *Gal4* (*UAS-NaChBac* or *UAS-Kir2.1*) (Baines et al., 2001; White et al., 2001; Cole et al., 2005; Nitabach et al., 2005). The *UAS-NaChBac* transgene is derived from a gene encoding a bacterial  $\text{Na}^+$  channel, which has the characteristics of high open probability and low inactivation, thus driving membrane voltage to a more depolarized and easily excited state. Expression of the  $\text{Na}^+$  channel in *Tdc2*-positive cells resulted in a decrease in sleep of 56.5%, corresponding to a loss of  $\sim 346$  min (Fig. 4A,B). The loss of sleep was specific to the nighttime, with no significant sleep loss during the daytime, which may be indicative of normally high octopamine activity during the day. This hypothesis is consistent with mammalian studies in which the noradrenergic cells of the locus ceruleus fire primarily during the active period (Aston-Jones and Bloom, 1981). We also found a corresponding increase in sleep latency in these flies. They took, on average, 74 min longer to fall asleep after lights off (Fig. 4B). Flies expressing *NaChBac* in *Tdc2*-positive cells also showed a decreased arousal threshold, suggesting that they are easily awakened during the night (Fig. 4C). Bout analysis indicated that nighttime sleep bouts were shorter in duration (supplemental Table 2, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). In addition, there was a significant increase in daytime bout number (supplemental Table 2, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material), which may reflect increased homeostatic drive resulting from the reduced sleep at night. However, the animals were unable to maintain long sleep bouts even under these conditions.

We also expressed a hyperpolarizing  $\text{K}^+$  channel transgene under the control of the *Tdc2-Gal4* driver and found that this produced an increase in total sleep (Fig. 5A,B). *Kir2.1* is an in-



**Figure 4.** Baseline sleep phenotype produced by depolarizing *Tdc2*-positive neurons. Female flies carrying a *UAS-NaChBac* transgene under the control of *TDC2-Gal4* were assayed for sleep. **A**, Six days of baseline sleep recording of eight animals each. *w;Tdc2-Gal4/NaChBac* + flies (gray dashed line) show significantly less sleep than *Iso31* controls (black solid line). Dark bars and white bars indicate nighttime and daytime, respectively. CT, Circadian time; ZT, Zeitgeber time. **B**, Nighttime sleep is significantly decreased in *w;Tdc2-Gal4/NaChBac* + flies (mean ± SEM; *w;Tdc2-Gal4/NaChBac* +, 267 ± 31, n = 33; *Iso31*, 613 ± 9, n = 58;  $p \leq 0.0001$ , two-way ANOVA). The latency to sleep is significantly longer in *w;Tdc2-Gal4/NaChBac* + flies (mean ± SEM; *w;Tdc2-Gal4/NaChBac* +, 107 ± 20, n = 33; *Iso31*, 34 ± 3, n = 58;  $p \leq 0.01$ , two-way ANOVA). **C**, Arousal threshold in the *w;Tdc2-Gal4/NaChBac* + flies. The animals were given three levels of stimulation to determine whether they were arousable. All the *w;Tdc2-Gal4/NaChBac* + flies were more arousable at each stimulation level, and there was a much lower percentage of animals that did not respond at all (*w;Tdc2-Gal4/NaChBac* +, n = 33; *Iso31*, n = 58).

ward rectifying  $K^+$  channel that has a high open probability and no inactivation. Expression of this channel hyperpolarizes neurons and decreases membrane resistance, thus making it more difficult for membrane potential to reach threshold for firing action potentials (Baines et al., 2001). Consistent with a previous report, we found that locomotor activity decreased when *Tdc2-Gal4* was used to express the inward rectifying  $K^+$  channel (*UAS-Kir2.1*) (data not shown). Analysis of sleep parameters, however, revealed that it was actually an increase in sleep that accounted for the phenotype (Fig. 5A,B). Flies expressing *UAS-Kir2.1* in *Tdc2* cells showed, on average, a 174 min increase in sleep and also displayed changes in several sleep measures such as latency and arousal threshold. These flies also showed a decrease in latency to sleep (Fig. 5B) and an increased arousal threshold during sleep, requiring more stimulation to wake up compared with controls (Fig. 5C). In addition, they showed a trend toward longer bouts of sleep during the night, although the major increase in sleep came from the increased number of sleep bouts during the day (supplemental Table 2, available at [www.jneurosci.org](http://www.jneurosci.org) as supplement-



**Figure 5.** Baseline sleep phenotype produced by hyperpolarizing *Tdc2*-positive neurons. Female flies carrying a *UAS-Kir2.1* transgene under the control of *Tdc2-Gal4* flies were assayed for sleep. **A**, Four days of baseline sleep recording of 16 animals for each group. *w;Tdc2-Gal4/Kir2.1;RC1* flies (gray dashed line) show significantly more sleep than their controls, *w;RC1;RC1* (black solid line). Dark bars and white bars indicate nighttime and daytime, respectively. CT, Circadian time; ZT, Zeitgeber time. **B**, Total sleep is significantly increased in *w;Tdc2-Gal4/Kir2.1;RC1* flies (mean ± SEM; *w;Tdc2-Gal4/Kir2.1;RC1*, 829.19 ± 14.14, n = 47; *w;RC1;RC1*, 655.35 ± 15.30, n = 47;  $p \leq 0.0001$ , two-way ANOVA). The latency to sleep is significantly lower in *Tdc2* × *Kir2.1* flies (mean ± SEM; *w;Tdc2-Gal4/Kir2.1;RC1*, 13.86 ± 1.19, n = 47; *w;RC1;RC1*, 45.68 ± 8.40, n = 47;  $p \leq 0.001$ , two-way ANOVA). **C**, Arousal threshold in *w;Tdc2-Gal4/Kir2.1;RC1* flies. The animals were given three levels of stimulation to determine whether they were arousable. All animals that responded to the first stimulation were aroused on stronger stimulation. Compared with controls, an increased percentage of *w;Tdc2-Gal4/Kir2.1;RC1* flies did not respond to any of the three levels of stimulation. At the strongest stimulation, more control flies responded than *w;Tdc2-Gal4/Kir2.1;RC1* flies (*w;Tdc2-Gal4/Kir2.1;RC1*, n = 47; *w;RC1;RC1*, n = 47).

tal material). The relative lack of an increase in nighttime sleep length may result from a ceiling effect of sleep at night.

Although sleep levels and architecture are clearly affected in flies expressing sodium or potassium channels in *Tdc2* neurons, it is possible that altered activity levels contribute to the overall phenotype. Thus, we also examined peak activity levels and activity while awake. The *Tdc2-Gal4* females expressing *NaChBac* showed significantly lower peak activity (supplemental Table 1, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material) as well as decreased activity while awake compared with the outcrossed background control (mean ± SEM; *w;Tdc2-Gal4/NaChBac* +, 1.7 ± 0.06; *Iso31*, 2.09 ± 0.06;  $p \leq 0.01$ , Kruskal–Wallis test). However, it is unlikely that a decrease in waking activity underlies a reduced sleep phenotype. We also found significantly reduced waking activity in animals expressing the  $K^+$  channel under the control of the *Tdc2* driver (supplemental Table 1, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). This decrease in

waking activity is similar to the decreased waking activity seen with the *Tdc2*<sup>RO54</sup> mutant, in which levels of both octopamine and tyramine are low (mean  $\pm$  SEM; *w;Tdc2-Gal4/Kir2.1;RC1*,  $2.14 \pm 0.06$ ; *w;RC1;RC1*,  $3.87 \pm 0.97$ ;  $p \leq 0.0001$ , Kruskal–Wallis test). Thus, reducing electrical activity in octopamine-producing cells has the same effect on activity and sleep as a mutation (*Tdc2*<sup>RO54</sup>) that decreases levels of octopamine.

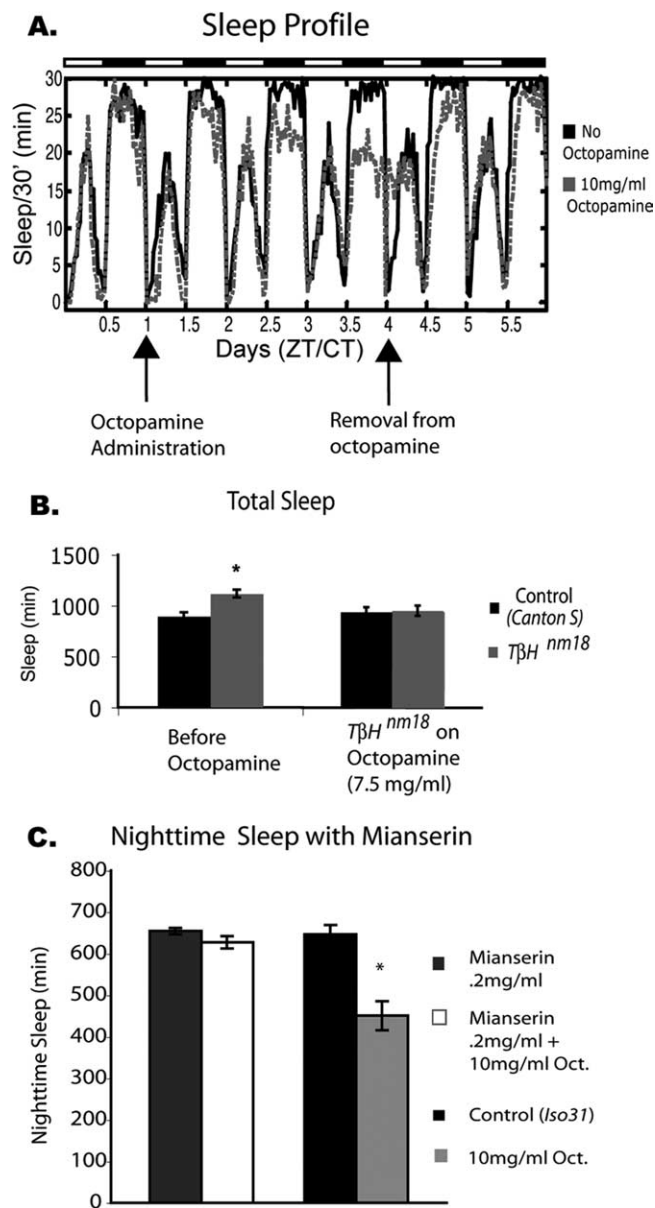
To ensure that the effect on sleep caused by loss of octopamine signaling is specific to neuronal *Tdc2* and not to a global loss of *Tdc*, we made use of a *Tdc1-Gal4* driver that is expressed in non-neuronal cells. Expression of the *UAS-B16B* transgene (*NaChBac* channel) under the control of this driver produced no significant change in sleep (supplemental Fig. 1, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). Thus, we conclude that the sleep phenotype observed in the *Tdc2*<sup>RO54</sup> mutant or produced by manipulations of *Tdc* cells is specific to the neuronal form of *Tdc*, *Tdc2*. Unfortunately, the *UAS-Kir2.1* channel proved to be lethal with *Tdc1-Gal4*.

Given that the *NaChBac* channel increases excitability of *Tdc2* cells and thereby presumably stimulates release of octopamine/tyramine, we asked whether overexpressing *Tdc2* would have the same effect. As predicted, we found that overexpression of *Tdc2* in *Tdc2*-producing cells resulted in a 300 min decrease in nighttime sleep (supplemental Fig. 2, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). As with the *NaChBac* channel, this appears to be a nighttime-specific sleep loss. There was also a decrease in sleep with the overexpression of *Tdc1* in *Tdc2*-producing cells (data not shown).

#### Oral administration of octopamine reduces sleep in flies

Because the *Tdc2*<sup>RO54</sup> and *TβH*<sup>nm18</sup> mutants change levels of octopamine and tyramine, it is important to dissociate the effects of the two to identify the transmitter responsible for the sleep phenotype. As noted previously, *Tdc2*<sup>RO54</sup> decreases levels of both tyramine and octopamine, whereas the *TβH*<sup>nm18</sup> decreases octopamine but increases tyramine. To determine whether a change in octopamine is sufficient to regulate sleep, we placed a wild-type isogenic line, *Iso31*, on octopamine-containing food for 3 d. It was shown previously that animals fed 10 mg/ml octopamine have increased levels of this neurotransmitter, particularly in the brain (Barron et al., 2007). Supporting this finding, ingested octopamine rescues the egg-laying phenotype displayed by *TβH*<sup>nm18</sup> and *Tdc2*<sup>RO54</sup> mutants (Monastirioti et al., 1996; McClung and Hirsh, 1999; Cole et al., 2005). We found that flies fed 10 mg/ml octopamine had ~200 min less nighttime sleep than control flies maintained on sucrose-agar alone (Fig. 6A) (dose–response curve is shown in supplemental Fig. 3A, available at [www.jneurosci.org](http://www.jneurosci.org) as supplemental material). Thus, similar to the *UAS-NaChBac* effect, this was a nighttime-specific effect. After the removal of octopamine, these flies showed a corresponding sleep rebound of ~70 min (mean  $\pm$  SEM; control,  $289.90 \pm 13.92$ ,  $n = 40$ ; 10 mg/ml octopamine,  $367.53 \pm 16.91$ ;  $n = 40$ ;  $p \leq 0.001$ , two-way ANOVA). In addition to the effect it had on wild-type flies, a lower concentration of orally administered octopamine (7.5 mg/ml octopamine) was able to restore the sleep phenotype of the *TβH*<sup>nm18</sup> mutant to control levels (Fig. 6B). This concentration of octopamine produced no significant change in sleep in the *Canton S* strain, which is the background of the *TβH*<sup>nm18</sup> mutants.

To exclude the possibility that the sleep phenotype was attributable to some nonspecific toxicity associated with octopamine, we attempted to block the effect by inhibiting octopamine signaling. Thus, we coadministered 0.2 mg/ml mianserin, which acts by



**Figure 6.** Oral administration of octopamine decreases sleep in *Iso31* flies and *TβH*<sup>nm18</sup> mutant flies. Female flies of the *Iso31* strain were treated with octopamine and assayed for sleep. **A**, The sleep profile shows 1 d of baseline sleep, followed by administration of 10 mg/ml octopamine (gray dashed line) for 3 d. Octopamine was then removed and sleep was assayed for 2 more days. Control flies (black line) were fed normal food during this time. Arrows indicate the time when food was changed. The data are from 16 animals for each group. There was a significant decrease in nighttime sleep for the *Iso31* flies on 10 mg/ml octopamine (mean  $\pm$  SEM; control,  $612.04 \pm 8.47$ ,  $n = 40$ ; 10 mg/ml octopamine,  $409.33 \pm 19.77$ ,  $n = 40$ ;  $p \leq 0.0001$ , two-way ANOVA). CT, Circadian time; ZT, Zeitgeber time. **B**, Sleep in *TβH*<sup>nm18</sup> mutant flies is restored to control levels through the administration of octopamine. Before octopamine administration, *TβH*<sup>nm18</sup> flies showed a significant increase in total sleep time; after administration of octopamine, sleep in *TβH*<sup>nm18</sup> flies was not significantly different from that of control (*Canton S*) flies not on octopamine (*Canton S*,  $796 \pm 55$  min; *Canton S* plus 7.5 mg/ml octopamine,  $792 \pm 54$  min;  $n = 16$  for each group). **C**, Coadministration of mianserin blocks the effect of octopamine on nighttime sleep. When 10 mg/ml octopamine was administered along with 0.2 mg/ml mianserin, there was no longer a significant drop in nighttime sleep (compare with columns on the right). Sleep in flies treated with mianserin alone was not significantly different from that of controls [mean  $\pm$  SEM; mianserin,  $654 \pm 8$ ,  $n = 32$ ; mianserin plus 10 mg/ml octopamine,  $627 \pm 15$ ,  $n = 32$ ; *Iso31* (Control),  $647 \pm 22$ ,  $n = 32$ ; 10 mg/ml octopamine,  $450 \pm 35$ ,  $n = 32$ ;  $p \leq 0.001$ , two-way ANOVA].



inhibiting octopamine-induced cAMP increase (Fig. 6C) (Maqueira et al., 2005). Coadministration of mianserin almost completely blocked the effect of feeding 10 mg/ml octopamine to flies. This demonstrates that the loss in sleep produced by octopamine feeding is attributable to the ingestion of octopamine itself and not attributable to toxic, nonphysiological effects. In addition, these data suggest that the effects of octopamine on sleep are mediated by  $\beta$  receptors.

Wild-type flies fed tyramine did not show significant changes in sleep amount (supplemental Fig. 3B, available at www.jneurosci.org as supplemental material). In addition, neither octopamine nor tyramine produced a significant change in activity while awake (supplemental Fig. 3C, available at www.jneurosci.org as supplemental material).

### Octopamine acts via neuronal PKA to decrease sleep

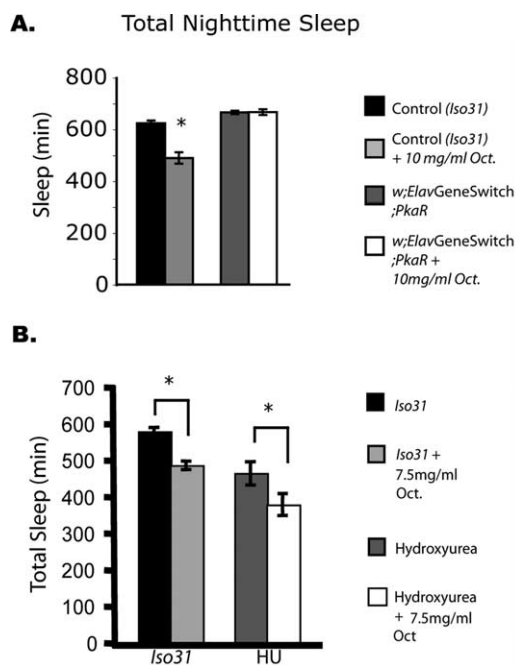
cAMP-dependent protein kinase (PKA) signaling plays an important role in sleep in *Drosophila* (Hendricks et al., 2001; Joiner et al., 2006). It is also known to be coupled to some of the octopamine and tyramine G-protein-coupled receptors (Evans and Maqueira, 2005). Indeed, the blocker of octopamine used above inhibits cAMP signaling. To directly address whether the effect of octopamine on sleep is through PKA-dependent pathways, we expressed the regulatory subunit of PKA (*PKAr*), which inhibits activity of PKA, under the control of an *Elav*GeneSwitch driver (Osterwalder et al., 2001; Joiner et al., 2006). The use of the *Elav*-GeneSwitch driver allowed us to inducibly express the regulatory subunit in all neurons. We found that expression of *PKAr* in adult neuronal tissue rendered the flies insensitive to the sleep-reducing effects of octopamine (Fig. 7A), supporting the idea that octopamine is acting through a PKA-dependent pathway to promote arousal.

Because the MB is implicated in the effects of PKA on sleep (Joiner et al., 2006), we sought to determine whether this is also the site of octopamine action on sleep. Thus, we ablated the mushroom bodies with hydroxyurea as described previously (de Belle and Heisenberg, 1994) and then treated these flies with octopamine. The sensitivity to octopamine was intact despite the absence of the MBs (Fig. 7B). As expected, the MB ablation itself reduced sleep, but there was an additional decrease produced by feeding octopamine (Fig. 7B). The decrease was 87 min, which was comparable with the amount of sleep lost when *Iso31* flies were fed 7.5 mg/ml octopamine (82 min). To verify that the MBs were ablated, we followed up the behavioral analysis of all flies with anatomical analysis of the brain. All flies that still contained some  $\alpha$  and  $\beta$  lobes were eliminated from analysis.

We also examined the effects of octopamine on flies expressing *PKAr* under control of an inducible mushroom body GeneSwitch driver (*MBSwitch*) and found that these flies were still sensitive to octopamine (data not shown). Our inability to block effects of octopamine on sleep by inhibiting PKA signaling in MBs supports the finding that elimination of MBs by hydroxyurea does not block the wake-promoting effects of octopamine.

## Discussion

Biogenic amines play many important roles in mammals, with several having significant effects on sleep–wake states. Thus, dopamine, serotonin, and norepinephrine are all important for regulating states of arousal. We hypothesized that, like its counterparts in mammals, the invertebrate neurotransmitter octopamine would be important for arousal in *Drosophila*. That prediction was supported by the data reported here. We find that decreases in levels of octopamine increase sleep, whereas increas-



**Figure 7.** Effects of octopamine are mediated by PKA and are independent of the mushroom body. **A**, Total nighttime sleep in control (*Iso31*) and *Elav*GeneSwitch flies crossed with *UAS-BDK33* (*PKAr*). The control on 7.5 mg/ml octopamine shows a significant decrease in nighttime sleep, whereas flies expressing *PKAr* under the control of *Elav*GeneSwitch do not show a decrease in sleep in response to octopamine [mean  $\pm$  SEM; Control (*Iso31*), 623  $\pm$  11,  $n$  = 24; control plus octopamine, 489  $\pm$  22,  $n$  = 24;  $p$   $\leq$  0.001, two-way ANOVA; *Elav*GeneSwitch, 665  $\pm$  7,  $n$  = 51; *Elav*GeneSwitch plus octopamine, 667  $\pm$  11,  $n$  = 51]. **B**, *Iso31* flies subjected to hydroxyurea treatment to ablate the mushroom body still show sensitivity to the sleep-reducing effects of octopamine. Hydroxyurea-treated flies show a significant decrease in sleep compared with controls; they also show an additional decrease in sleep when placed on 7.5 mg/ml octopamine (mean  $\pm$  SEM; control, 583  $\pm$  11,  $n$  = 24; control plus 7.5 mg/ml octopamine, 491  $\pm$  12,  $n$  = 24;  $p$   $\leq$  0.01, two-way ANOVA; HU, 469  $\pm$  32,  $n$  = 24; HU plus 7.5 mg/ml octopamine, 382  $\pm$  30,  $n$  = 22;  $p$   $\leq$  0.01, two-way ANOVA).

ing octopamine causes a decrease in sleep. In addition, although the mammalian data have been contradictory with respect to the role norepinephrine plays in total sleep, we find that octopamine decreases total sleep time. The mammalian data are complicated in part because perturbations of the norepinephrine pathway result in changes in the levels of dopamine (Ouyang et al., 2004). The use of *Drosophila* allows us to examine specifically the role of octopamine without perturbing dopamine signaling.

By modulating the excitability of octopamine-producing cells, we were able to manipulate the output of these cells. In mammals, one can record from specific cell populations to determine when the cells fire action potentials. Although this assay is difficult to do in flies, we were able to electrically modulate the cells through expression of  $K^+$  and  $Na^+$  ion channels. We found that, when octopamine-producing cells were more depolarized (expression of a  $Na^+$  channel), the animal was awake more and unable to stay asleep, whereas when the cells were hyperpolarized (expression of a  $K^+$  channel), the animals slept more.

Based primarily on larval crawling assays, octopamine and tyramine were implicated previously in locomotor behavior (O'Dell, 1994; Gong et al., 2004; Saraswati et al., 2004; Scholz, 2005). Specifically, larvae move slower through quadrants when they have decreased octopamine levels (the *TBH<sup>nm18</sup>* and *Tdc2<sup>RO54</sup>* mutants). More recent work showed that adult *Tdc2<sup>RO54</sup>* flies also have a decrease in locomotor activity attributable to the lack of tyramine (Hardie et al., 2007). Our data show-

ing differences in activity in the *Tdc2*<sup>RO54</sup> and the *TβH*<sup>nm18</sup> mutants support the claim that tyramine plays an important role in locomotion. Thus, whereas increased levels of tyramine in *Tbh* mutants increase activity, decreased levels in *Tdc* mutants decrease locomotor activity. However, both mutations increase sleep, which is most likely attributable to the loss of octopamine. In addition to overall sleep, we find that other sleep parameters such as latency to sleep and arousal threshold are affected in flies carrying these mutations. We infer that tyramine plays a role in locomotion, but octopamine specifically affects arousal states.

Studies of other invertebrate species support a role for octopamine in arousal (Corbet, 1991). In fact, octopamine agonists are potential natural pesticides because they cause insect species to “walk off” the leaves (Roeder, 1999). As in *Drosophila*, changes in octopamine levels affect behavior in honey bees, as demonstrated through feeding and injection of octopamine as well as through analysis of endogenous levels of octopamine. Fussnecker et al. (2006) showed that injections of octopamine promote flying in honeybees. In addition, octopamine and tyramine regulate other behaviors in honeybees such as hive maintenance and foraging (Schulz and Robinson, 1999, 2001; Wagener-Hulme et al., 1999; Barron et al., 2002). Octopamine and tyramine also modulate sensory input in honeybees (Kloppenborg and Erber, 1995; Scheiner et al., 2002). In the locust, octopamine mediates heightened arousal in response to new visual stimuli (Bacon et al., 1995). Bacon et al. found that a specific subset of octopamine-producing neurons in the brain of the locust fires during the presentation of new visual stimuli, causing dishabituation of the descending contralateral movement detector interneuron. Interestingly, application of endogenous octopamine can mimic this state of heightened arousal. Our study suggests that octopamine serves to promote arousal in *Drosophila*. It is possible that the increased arousal we see with too much octopamine, or decreased arousal with too little, is a result of improper gating of sensory stimuli, but without electrophysiological data we are unable to draw any conclusions. Note also that the *Tdc2* cells important for sleep and arousal in the fly brain have not been identified yet.

In previous studies, octopamine was fed to flies to rescue or verify a phenotype of the *TβH*<sup>nm18</sup> flies. The ability of octopamine to rescue egg laying in *TβH*<sup>nm18</sup> mutants was assayed in this manner, because *TβH*<sup>nm18</sup> flies are unable to release eggs. Animals were placed on different levels of octopamine, and 10 mg/ml octopamine over a period of 6 d provided maximal rescue (Monastirioti et al., 1996). Using the same concentration, we found that a steady increase in octopamine levels led to a decrease in nighttime sleep. Based on the specific effect on nighttime sleep, we speculate that octopamine levels are already high during the daytime, thereby precluding any effects of an increase. This analysis is supported by the Na<sup>+</sup> channel data in which a significant decrease in total sleep was found only during the nighttime sleep periods. We speculate that, normally, activity of these cells is low at night, and so expression of the Na<sup>+</sup> channel causes them to fire more and release octopamine at an abnormal time, thereby producing a decrease in sleep. Similar results, indicating nighttime-specific effects, were obtained with overexpression of *Tdc2*. Work in other insects also supports the idea of modulated octopamine release. Pribbenow and Erber (1996) demonstrated that honeybees who are already in a heightened arousal state of antennae scanning do not change scanning frequency in response to octopamine administration, but, in animals scanning at a low frequency, injections of octopamine significantly increase scanning (Pribbenow and Erber, 1996).

Our data suggest that the effects of octopamine are mediated

through PKA-dependent signaling. In mammals, there are nine different adrenergic receptors, some of which signal through PKA (Hoffman and Lefkowitz, 1996). The α1 adrenergic receptor is the only receptor associated with a wake-promoting effect in that the agonist methoxamine causes an increase in waking (Hilakivi and Leppävuori, 1984; Monti et al., 1988). However, the antagonist has no effect on total sleep (Benington et al., 1995; Berridge and España, 2000). It is important to note that the α1 receptor in mammals is thought to be coupled to phospholipase C and G<sub>q</sub> (Ramos and Arnsten, 2007). The β adrenergic receptors (which are coupled to cAMP and PKA) probably do not have specific effects on sleep in mammals because, contrary to known effects of norepinephrine, the agonist increases sleep and the antagonist decreases sleep (Monti et al., 1988). Studies in *Drosophila* may be better able to identify biogenic amine receptors relevant for sleep because of the ease of genetic manipulation. Many G-protein-coupled receptors in *Drosophila* display activity that allows their bona fide classification as octopamine receptors (Evans and Maqueira, 2005). Our data here suggest that receptors sensitive to mianserin are likely to be involved in regulating fly sleep. Because mianserin inhibits cAMP signaling, these data not only further support a role for PKA but also implicate β receptors in octopamine action. We note that none of these receptors is known to display a circadian cycling profile.

Given that PKA was shown previously to regulate sleep in *Drosophila*, we are starting to see a link between the various molecules that affect *Drosophila* sleep. Interestingly, however, octopamine does not appear to act through the MBs, a structure known to mediate effects of PKA on sleep and also to express a class of octopamine receptors. Because flies lacking MBs still have substantial amounts of sleep, it is clear that other parts of the fly brain can drive sleep. The current study shows that even PKA can affect sleep in regions outside the MB. Defining the site of action of sleep-regulating molecules such as octopamine should help to identify these other brain regions.

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