

Figure S1. Deletion of *Bmal1* from SCN^{VIP} neurons via genetic cross.

To quantify the efficiency and specificity of *Bmal1* deletion in *VIP^{cre/cre}::Bmal1^{fl/fl}* mice we generated triple cross *VIP-ires-Cre::Bmal1^{fl/fl}::R26-loxSTOPlox-L10-GFP* mice. Example confocal images of the SCN of *VIP^{cre/wt}::Bmal1^{wt/wt}::R26-loxSTOPlox-L10-GFP* (L10-GFP) mice showing 100% co-localization between VIP (L10-GFP, green) and *Bmal1* (red; left and top right). In *VIP^{cre/wt}::Bmal1^{fl/fl}::L10-GFP* mice, ~ 70% of VIP-positive neurons continued to express *Bmal1* (center right). In *VIP^{cre/cre}::Bmal1^{fl/fl}::L10-GFP* mice, 0% of VIP-positive neuron expresses *Bmal1* (bottom right), although *Bmal1* continued to be abundantly expressed in adjacent non-VIPergic neurons of the ventral SCN. White arrows indicate VIP-*Bmal1* co-localization, while blue arrows show absence of co-localization. Scale bars: left, 100 μ m and right, 20 μ m.

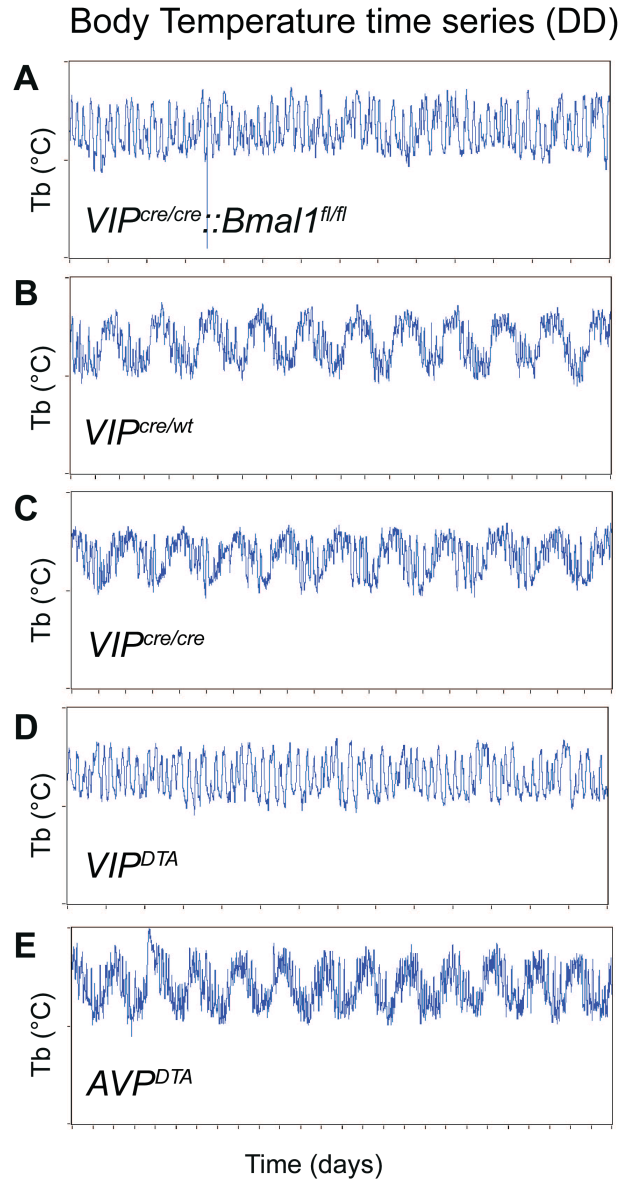


Figure S2. Tb rhythm waveforms in DD in $VIP^{cre/wt}$, $VIP^{cre/cre}$, $VIP^{cre/cre}::Bmal1^{fl/fl}$, VIP^{DTA} , and AVP^{DTA} mice. Related to Figure 1 and 3. Selective deletion of *Bmal1* in SCN^{VIP} neurons (A) has clear visible disruptive effective on the Tb waveform compared to $VIP^{cre/wt}$ (B) and $VIP^{cre/cre}$ (C) mice, yet these rhythms, per periodogram analysis, retain a circadian component in the $VIP^{cre/cre}::Bmal1^{fl/fl}$ mouse. Selective ablation of SCN^{VIP} neurons (D), but not SCN^{AVP} neurons (E), severely disrupts circadian rhythms of Tb.

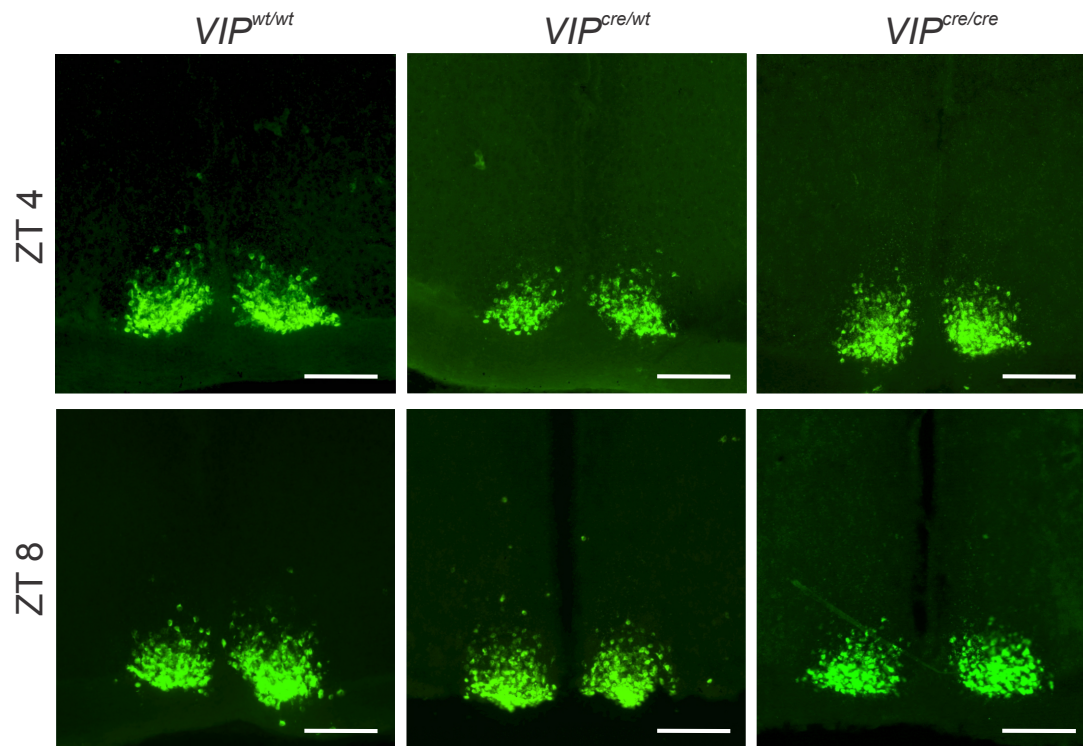


Figure S3. VIP mRNA expression in $VIP^{wt/wt}$, $VIP^{cre/wt}$ and $VIP^{cre/cre}$ conditions.

Related to Figure 1. RNAscope-based *in situ* hybridization for VIP mRNA within the SCN is shown from brains sacrificed at ZT4 (upper panels) and ZT8 (lower panels) for $VIP^{wt/wt}$, $VIP^{cre/wt}$ and $VIP^{cre/cre}$. VIP transcript levels are visually comparable across the wild type, heterozygous and homozygous condition in *VIP-ires-Cre* mice. Scale bars: 100 μm .

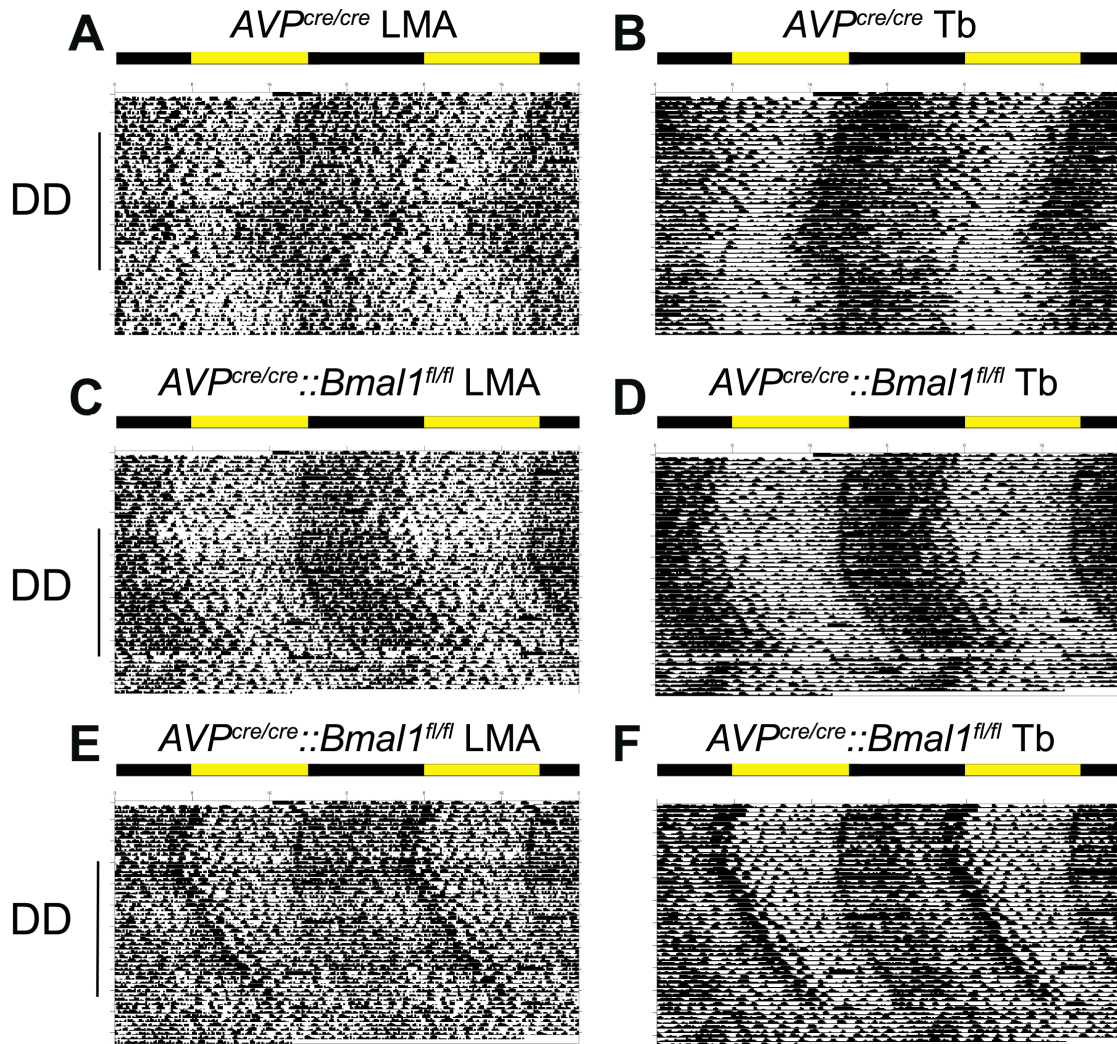


Figure S4. Loss of the molecular clock in SCN^{AVP} neurons on LMA and Tb rhythms.
Related to Figure 1. A-B. Example actogram (LMA; A,C,E) and Tb variation (B,D,F; grey scale: darker represents higher temperature) for an $AVP^{cre/cre}$ mouse (A-B) and two $AVP^{cre/cre}::Bmal1^{fl/fl}$ mice (C-F) during at least 10 days in light/dark condition (LD), followed by at least 3 weeks days in constant dark (DD), followed by at least seven days of LD. Note that $AVP^{cre/cre}::Bmal1^{fl/fl}$ mice retain coherent rhythms of LMA and Tb, but exhibit a lengthened period compared to $AVP^{cre/cre}$ mice.

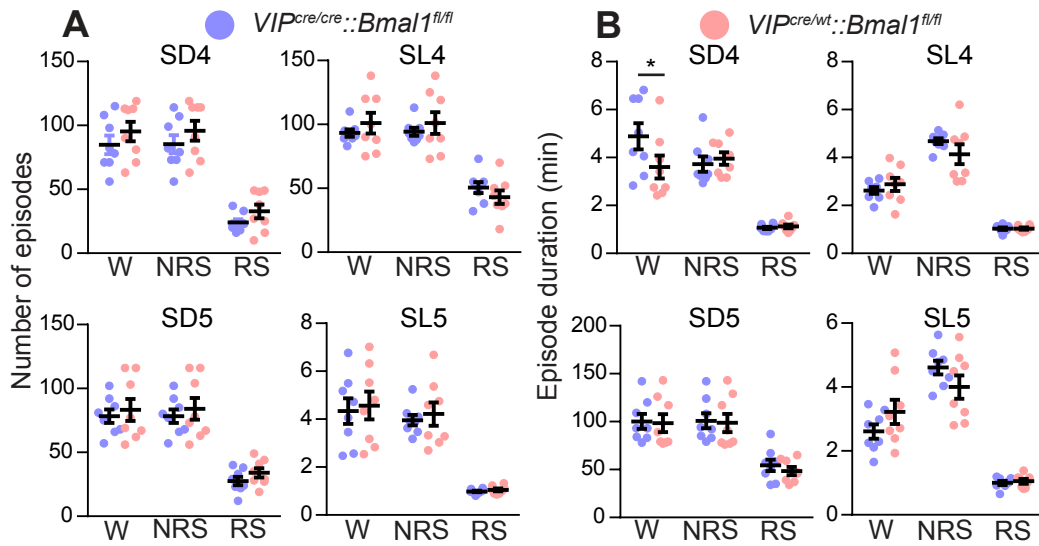


Figure S5. Sleep-wake qualitative changes following loss of the molecular clock in SCN^{VIP} neurons. Related to Figure 2. (A) Number of episodes (mean \pm s.e.m.) of wake (W), NREM sleep (NRS) or REM sleep (RS) and **(B)** episode duration (mean \pm s.e.m.) of W, NRS or RS during the 4th and 5th subjective dark (SD4, SD5), and 4th and 5th subjective light (SL4, SL5) periods in $VIP^{cre/cre}::Bmal1^{fl/fl}$ (blue, n = 8) and $VIP^{cre/wt}::Bmal1^{fl/fl}$ (red, n = 8) mice. * $p < 0.05$, two-way ANOVA followed by a *post hoc* Bonferroni test.

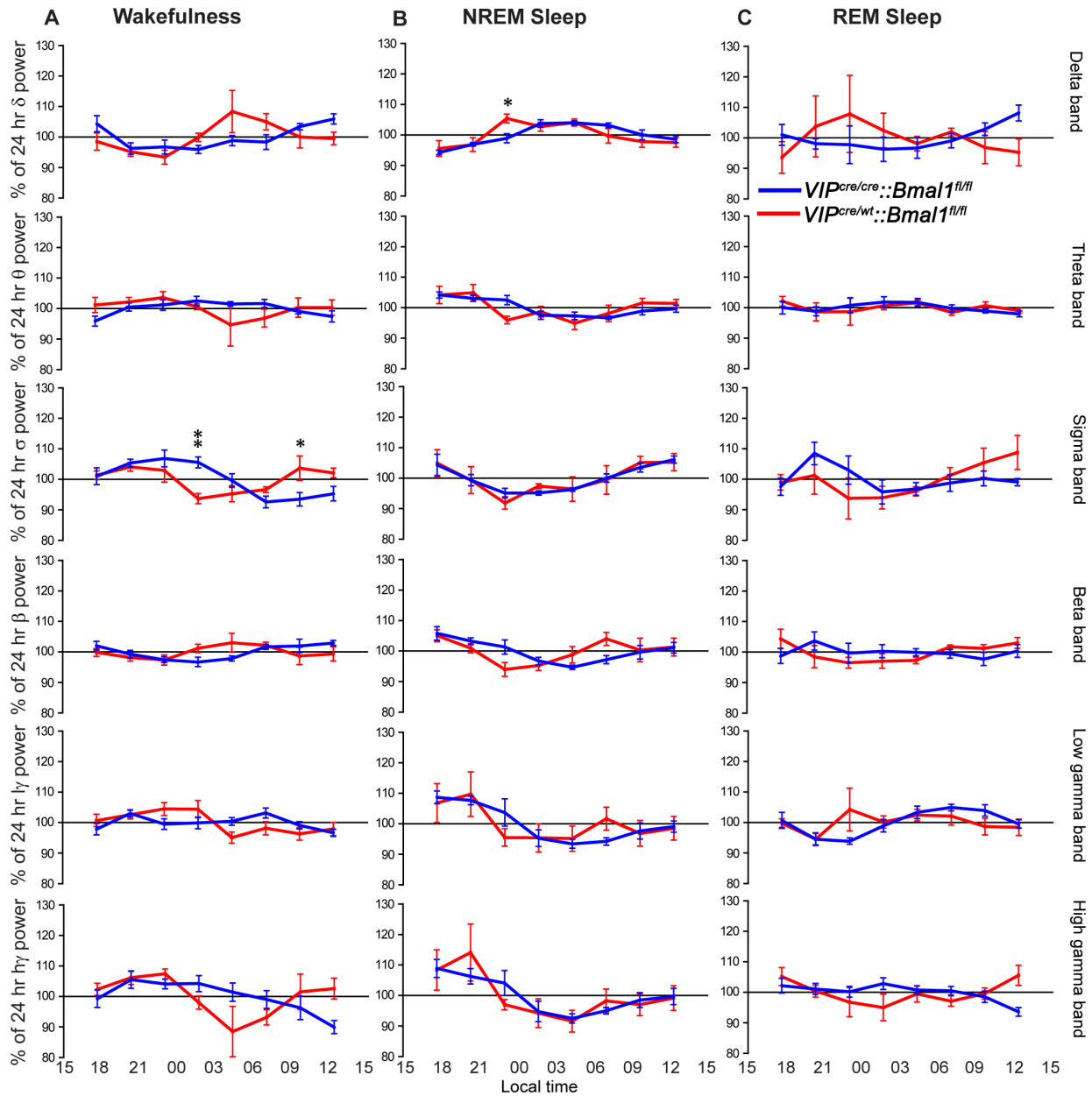


Figure S6. Sleep-wake cortical EEG power band activity across the 24 hr period following loss of the molecular clock in SCN^{VIP} neurons. Related to Figure 2. Delta (δ : 0.5–5 Hz), theta (θ : 5–9 Hz), sigma (σ : 9–15 Hz), beta (β : 15–30 Hz), low gamma (γ : 30–60 Hz) and high gamma (γ_H : 60–120 Hz) power bands (mean \pm s.e.m) across 24 hrs of the 4th day in constant darkness in *VIP^{cre/cre}::Bmal1^{fl/fl}* (blue, n = 6) and

$VIP^{cret/wt}::Bmal1^{fl/fl}$ (red, $n = 6$) mice. Power bands were averaged for each 3 hr period and expressed as a percentage of the 24 hr average power for each frequency band and each vigilance state. * $p < 0.05$, ** $p < 0.01$, two-way ANOVA followed by a *post hoc* Bonferroni test.

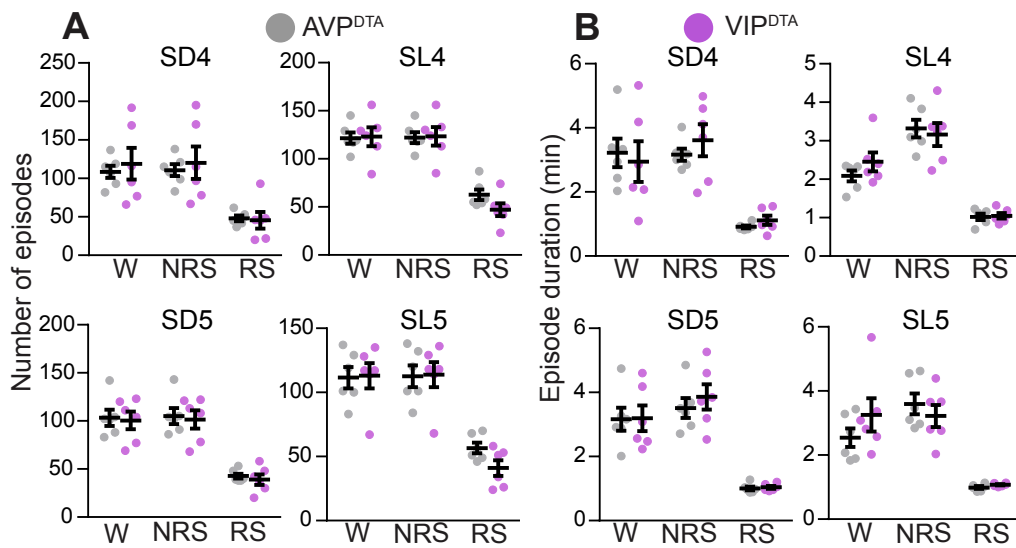


Figure S7. Sleep-wake qualitative changes following genetically-driven lesions of SCN^{VIP} or SCN^{AVP} neurons. Related to Figure 4. (A) Number of episodes (mean \pm s.e.m.) of wake (W), NREM sleep (NRS) or REM sleep (RS) and (B) episode duration (mean \pm s.e.m.) of W, NRS or RS during the 4th and 5th subjective dark (SD4, SD5), and 4th and 5th subjective light (SL4, SL5) periods in AVP^{DTA} (grey, $n = 6$) and VIP^{DTA} (purple, $n = 6$) mice. $p > 0.05$, two-way ANOVA followed by a *post hoc* Bonferroni's test.

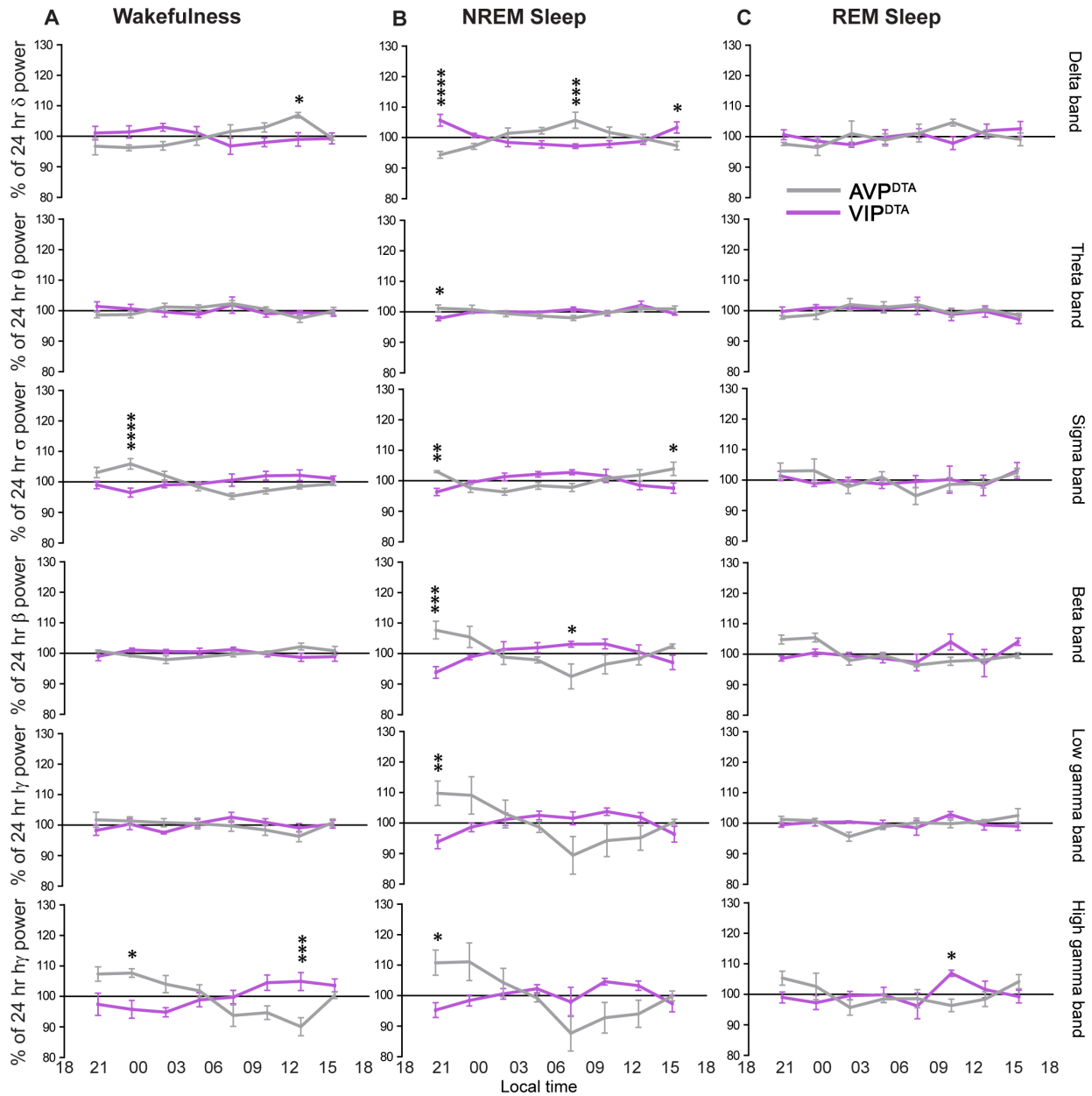
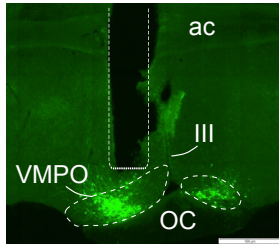
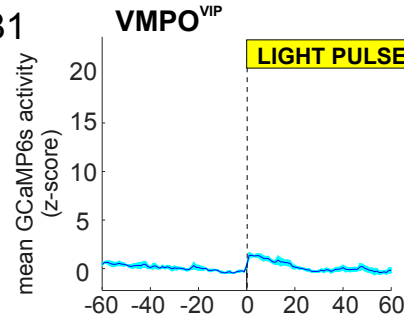
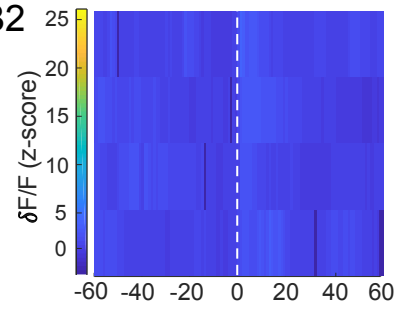
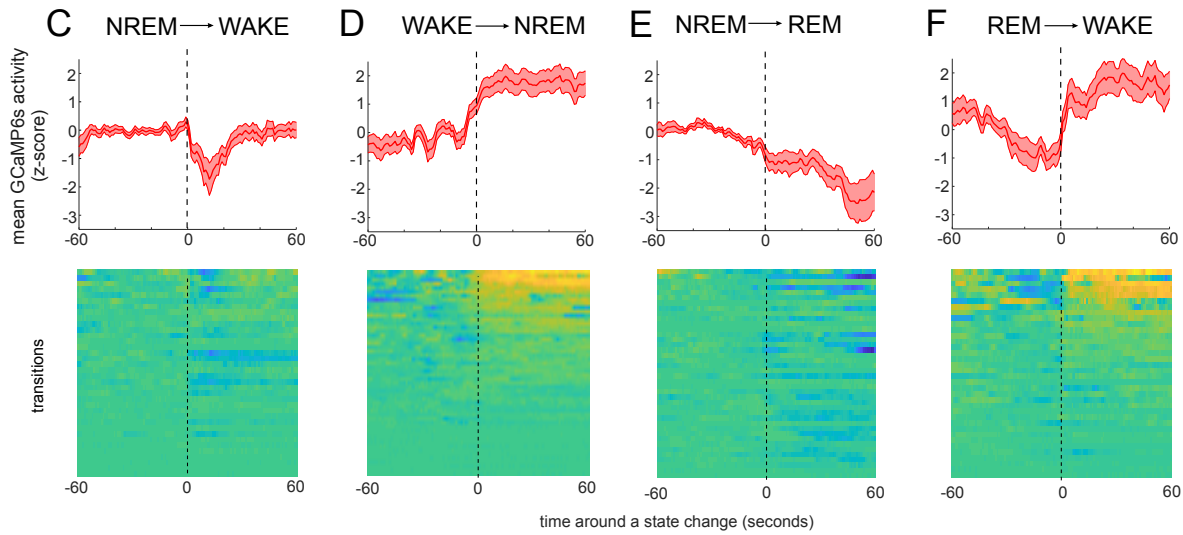


Figure S8. Sleep-wake cortical EEG power band activity across 24 hr period genetically-driven lesions of SCN^{VIP} or SCN^{AVP} neurons. Related to Figure 4. Delta (δ : 0.5–5 Hz), theta (θ : 5–9 Hz), sigma (σ : 9–15 Hz), beta (β : 15–30 Hz), low gamma (γ : 30–60 Hz) and high gamma (γ_H : 60–120 Hz) power band (mean \pm s.e.m.) activity across 24 hr of the 4th day in constant darkness in AVP^{DTA} (grey, n=5) and VIP^{DTA} (purple, n=6) mice. Power bands were averaged for each 3 hr period and expressed as a percentage

of the 24 hr average power for each frequency band and each vigilance state. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, **** $p < 0.0001$, two-way ANOVA followed by a *post hoc* Bonferroni test.

A**B1****B2**

SUBJECTIVE MORNING (CT 2-5)



SUBJECTIVE EVENING (CT 12-14)

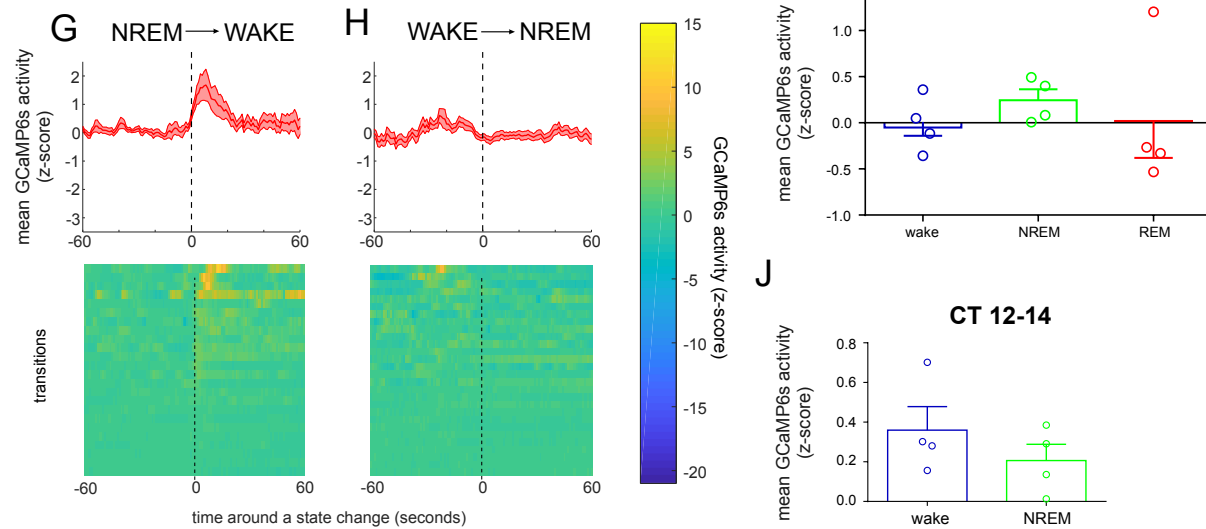


Figure S9. Activity in SCN^{VIP} neurons is behavioral state invariant. Related to Figure

7. (A) Photomicrograph depicting viral targeting to VIP-containing neurons in the ventromedial preoptic nucleus (VMPO^{VIP}) and placement of the photometry fiber. Abbreviations: III; 3rd ventricle, ac; anterior commissure, OC; optic chiasm **(B1)** Mean GCaMP6s activity in VMPO^{VIP} neurons over all mice at the initiation of the light pulse. Black dotted line indicates onset of the light pulse. Shaded area indicates s.e.m. **(B2)** Heatmap depicting GCaMP6s activity at the light pulse for each mouse. White dotted line indicates onset of the light pulse. **(C-J)** GCaMP6s activity at vigilance state transitions during the subjective morning (CT 2-5, C-F) and subjective evening (CT 12-14, G-H). Upper panels: means GCaMP6s activity across mice. Lower panels: Heatmaps depicting individual arousal state transitions from all mice. Paired t-tests (two-tailed) indicate no significant effect of the state transition upon GCaMP6s activity. **(I)** Mean (+ s.e.m) GCaMP6s activity during wake, NREM sleep and REM sleep between CT 2-5. RM one-way ANOVA indicated no significant effect of state upon GCaMP6s activity (Arousal state: $F(2, 6) = 0.2989, p = 0.7521$). **(J)** Mean (+ s.e.m) GCaMP6s activity during wake and NREM sleep between CT 12-14. Paired t-test (two-tailed) indicated no significant effect of state upon GCaMP6s activity ($p = 0.3128, n = 4$).

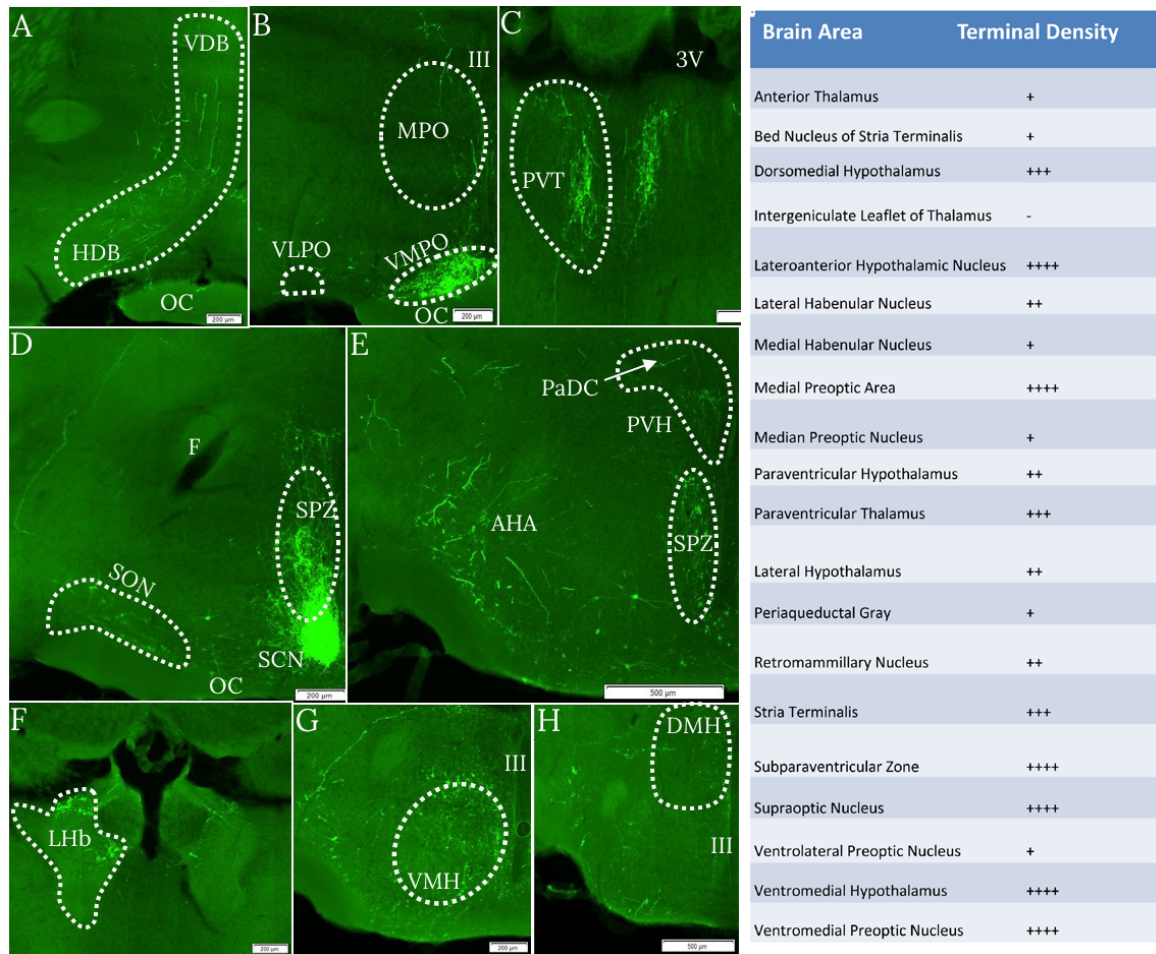


Figure S10. SCN^{VIP} neurons project densely to the subparaventricular zone and provide innervation to many other hypothalamic and non-hypothalamic structures.

Related to Figure 8. The anterograde tracer, AAV-FLEX-hrGFP was injected unilaterally into the SCN of *VIP-IRES-Cre* mice and axons filled with the tracer were assessed using a semi-quantitative method (see Methods) throughout the neuraxis. **(A-H)** Photomicrographs of hrGFP immunoreactivity (green) in the ventral diagonal band (VDB) and horizontal diagonal band (HDB) **(A)**, the ventromedial preoptic nucleus (VMPO) **(B)**, the paraventricular nucleus of the thalamus (PVT) **(C)**, the subparaventricular zone (SPZ) **(D)**, the paraventricular hypothalamus (PVH) and supraoptic nucleus (SON) **(E)**, the lateral habenula (LHb) **(F)**, the ventromedial hypothalamic nucleus (VMH) **(G)** and the

dorsomedial hypothalamic nucleus (DMH) (**H**). (**I**) Table showing density of axons in each brain region. Abbreviations: III, 3rd ventricle; AHA, anterior hypothalamic area, anterior part; DMH, dorsomedial hypothalamic nucleus; HDB, nucleus of the horizontal limb of the diagonal band; LHb, lateral habenular nucleus; MPO, medial preoptic nucleus; OC, optic chiasm; PaDC, paraventricular hypothalamic nucleus, dorsal cap; PVH, paraventricular hypothalamic nucleus; PVT, paraventricular thalamic nucleus; SON, supraoptic nucleus; SPZ, subparaventricular zone of the hypothalamus; VDB, nucleus of the vertical limb of the diagonal band; VLPO, ventrolateral preoptic nucleus; VMH, ventromedial hypothalamic nucleus; VMPO, ventromedial preoptic nucleus.

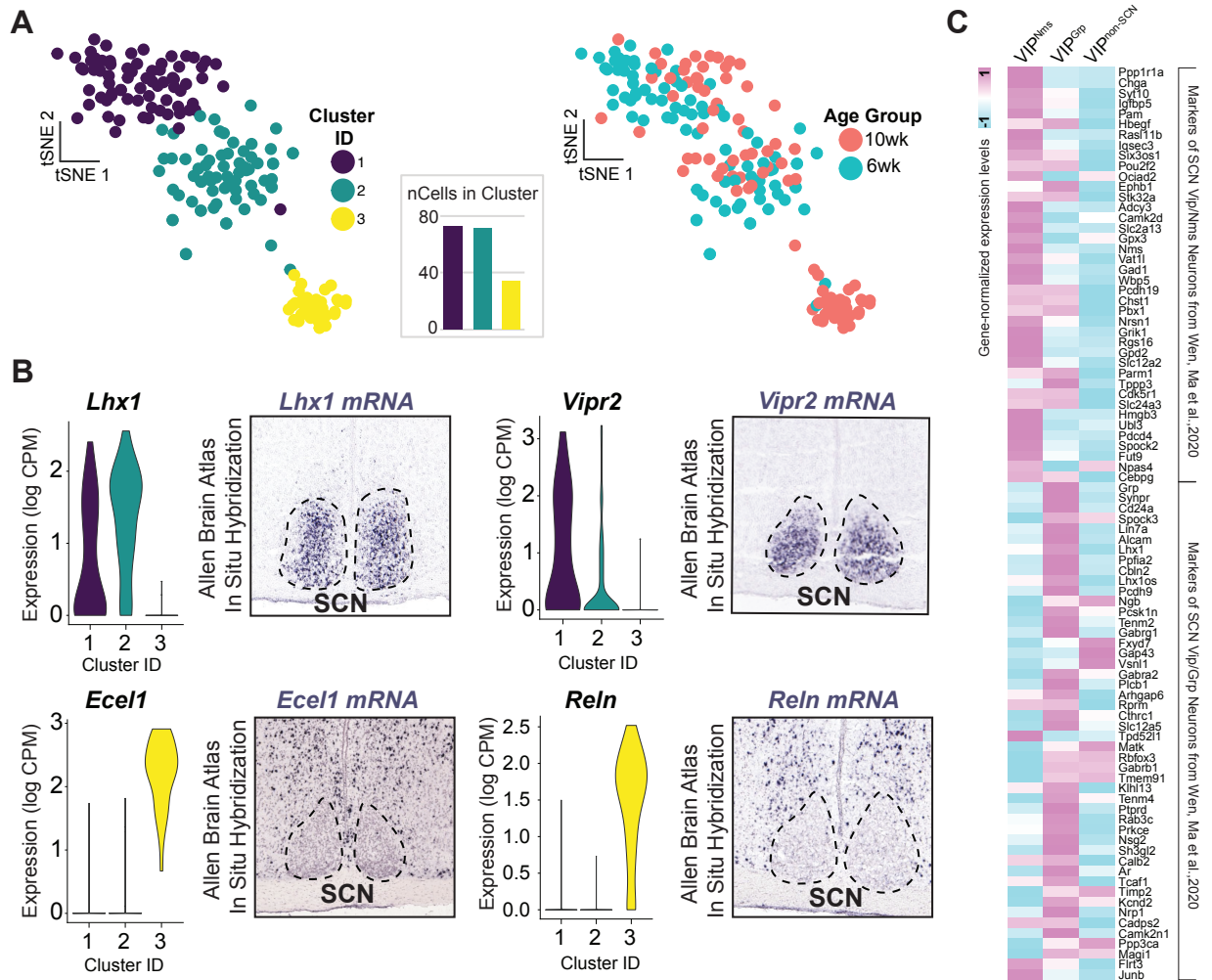


Figure S11. sNuc-seq analysis reveals three distinct VIP neuron populations, including two from the SCN. (A) t-Distributed Stochastic Neighbor Embedding (tSNE) plots of clustered VIP neurons, colored by cluster identity (left) or mouse age group (right). (B) Single-cell expression of genes enriched in or around the SCN, with corresponding in situ hybridization images from the Allen Mouse Brain Atlas: *Lhx1*, experiment 633; *Vipr2*, experiment 1104; *Ecel1*, experiment 70231305; and *Reln*, experiment 890. (C) Heatmap of cluster-level average expression (columns) of top marker genes (rows) for Vip/Grp neurons and Vip/Nms neurons identified by Wen, Ma et al., 2020⁴⁶.

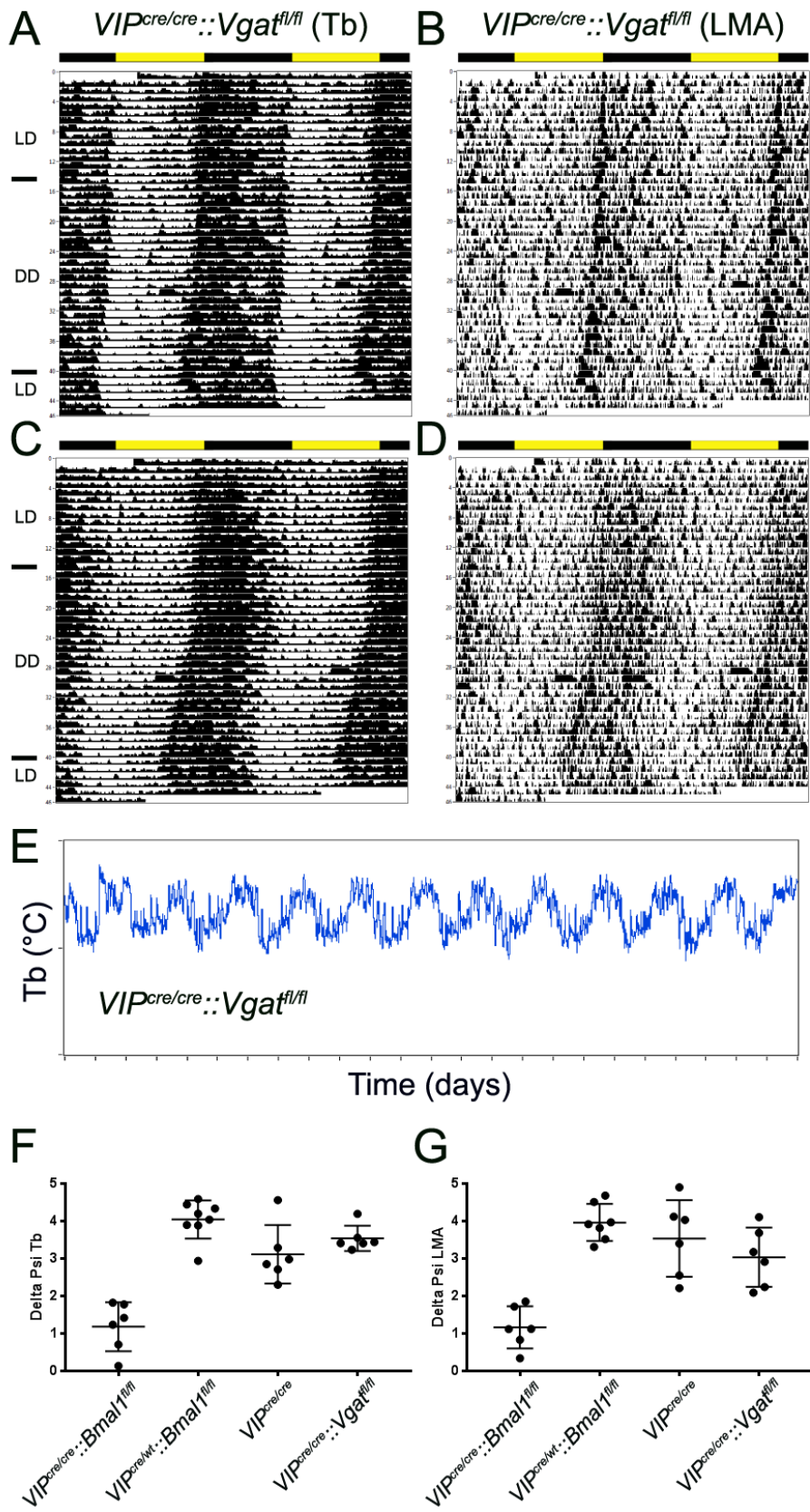
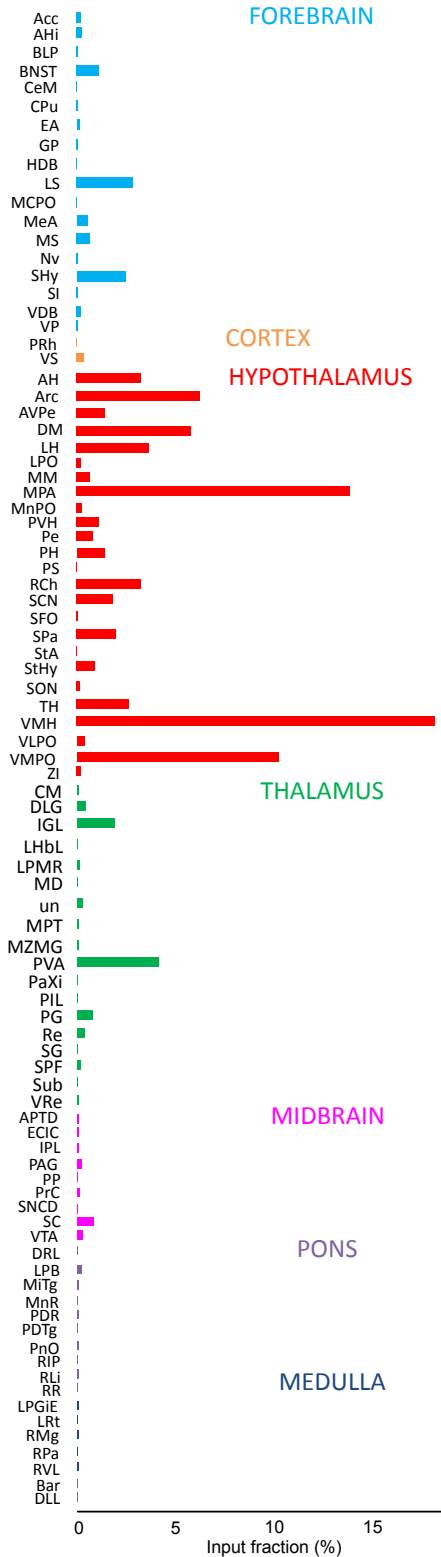


Fig S12. Genetic disruption of GABAergic transmission from SCN^{VIP} neurons. (A-D) Selective deletion of *Vgat* in SCN^{VIP} neurons produced no discernable disruption in the diurnal (LD) or circadian (DD) rhythms of Tb or LMA (n=2 *VIP^{cre/cre}::Vgat^{fl/fl}* mice shown). **(E)** High amplitude circadian rhythms of Tb were observed in the *VIP^{cre/cre}::Vgat^{fl/fl}* (cf. with *VIP^{cre/cre}::Bmal1^{fl/fl}* mouse in Fig S3A). **(F-G)** deletion of *Vgat* in SCN^{VIP} neurons reduced the variability of the phase angle of entrainment of Tb, but not LMA, as compared to *VIP^{cre/cre}* mice (i.e., putative hypomorphic condition). *VIP^{cre/cre}::Bmal1^{fl/fl}*, n=6; *VIP^{cre/wt}::Bmal1^{fl/fl}*, n=7; *VIP^{cre/cre}*, n=6; *VIP^{cre/cre}::Vgat^{fl/fl}*, n=6. Means \pm s.e.m.

Anatomical Abbreviations and Quantification for Figure 9



Abbreviation List: 3V, 3rd ventricle; ac, anterior commissure; Acc, accumbens nucleus; AH, anterior hypothalamic area, anterior part; Ahi, amygdalohippocampal area; alv, alveus of the hippocampus; APTD, anterior pretectal nucleus, dorsal part; Arc, arcuate hypothalamic nucleus; AVPe, anteroventral periventricular nucleus; Bar, Barrington's nucleus; BLP, basolateral amygdaloid nucleus, posterior part; BNST, bed nucleus of the stria terminalis; CeM, central amygdaloid nucleus, medial division; CM, central medial thalamic nucleus; CPu, caudate putamen (striatum); DLG, dorsal lateral geniculate nucleus; DLL, dorsal nucleus of the lateral lemniscus; DM, dorsomedial hypothalamic nucleus; DRL, dorsal raphe nucleus, lateral part; EA, extended amygdala; ECIC, external cortex of the inferior colliculus; fx, fornix; GP, globus pallidus; HDB, nucleus of the horizontal limb of the diagonal band; IGL, intergeniculate leaflet; IPL, interpeduncular nucleus, lateral subnucleus; LH, lateral hypothalamic area; LHbL, lateral habenular nucleus, lateral part; LPB, lateral parabrachial nucleus; LPGiE, lateral paragigantocellular nucleus, external part; LPMR, lateral posterior thalamic nucleus, mediorostral part; LPO, lateral preoptic area; LRt, lateral reticular nucleus; LSD, lateral septal nucleus, dorsal part; LSV, lateral septal nucleus, ventral part; LV, lateral ventricle; MCPO, magnocellular preoptic nucleus; MD, mediodorsal thalamic nucleus; MeA, medial amygdaloid nucleus, anterior part; MiTg, microcellular tegmental nucleus; MM, medial mammillary nucleus, medial part; MnPO, median preoptic nucleus; MnR, median raphe nucleus; MPA, medial preoptic area; MPT, medial pretectal nucleus; MS, medial septal nucleus; MZMG, marginal zone of the medial geniculate; Nv, navicular postolfactory nucleus; opt, optic tract; ox, optic chiasm; PAG, periaqueductal gray; PaXi, paraxiphoid nucleus of thalamus; PDR, posterodorsal raphe nucleus; PDTg,

posterodorsal tegmental nucleus; Pe, periventricular hypothalamic nucleus; PG, pregeniculate nucleus; PGMC, pregeniculate nucleus, magnocellular part; PGPC, pregeniculate nucleus, parvicellular part; PH, posterior hypothalamic nucleus; PIL, posterior intralaminar thalamic nucleus; PnO, pontine reticular nucleus, oral part; PP, peripeduncular nucleus; PrC, precommissural nucleus; PRh, perirhinal cortex; PS, parastrial nucleus; PVA, paraventricular thalamic nucleus, anterior part; PVH, paraventricular hypothalamic nucleus; PaDC, paraventricular hypothalamic nucleus, dorsal cap; Re, reuniens thalamic nucleus; RIP, raphe interpositus nucleus; RLi, rostral linear nucleus (midbrain); RMg, raphe magnus nucleus; RR, retrorubral nucleus; RVL, rostroventrolateral reticular nucleus; SC, superior colliculus; SCN, suprachiasmatic nucleus; SFO, subfornical organ; SG, suprageniculate thalamic nucleus; SHy, septohypothalamic nucleus; SHY, septohypothalamic nucleus; SI, substantia innominate; SNCD, substantia nigra, compact part, dorsal tier; SON, supraoptic nucleus; Spa, subparaventricular zone of the hypothalamus; SPF, subparafascicular thalamic nucleus; StA, strial part of the preoptic area; STLI, bed nucleus of the stria terminalis, lateral division; STLP, bed nucleus of the stria terminalis, lateral division, posterior part; STLV, bed nucleus of the stria terminalis, lateral division, ventral part; STMA, bed nucleus of the stria terminalis, medial division, anterior part; STMAL, bed nucleus of the stria terminalis, medial division, anterolateral part; Sub, submedial thalamic nucleus; Te, terete hypothalamic nucleus; un, unassigned thalamic area, located between the fasciculus retroflexus; VDB, nucleus of the vertical limb of the diagonal band; VLPO, ventrolateral preoptic nucleus; VMH, ventromedial hypothalamic nucleus; VMPO, ventromedial preoptic nucleus; VP, ventral pallidum; VRe, ventral reuniens thalamic nucleus; VS, ventral subiculum; VTA, ventral tegmental area, ZI, zona incerta.