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A wake-active locomotion circuit depolarizes a sleep-active neuron to switch on sleep

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Abbreviations: ArchT, archaerhodopsin from *Halorubrum* strain TP009; APTF-1, Activating enhancer binding Protein 2 Transcription Factor 1; ATR, all-trans-retinal; AVA, ventral cord interneuron class name; AVB, Ventral cord interneuron class name; AVD, ventral cord interneuron class name; AVE, ventral cord interneuron class name; AVJ, neuron class name; ASH, Amphid neuron class name; CEP, cephalic neuron class name; dFB, dorsal Fan-shaped Body; DIC, differential interference contrast; EEG, electroencephalogram; $\Delta F/F$, change of fluorescence over baseline; FLP-11, FMRF-Like Peptide 11; FLP-18, FMRF-Like Peptide 18; GCaMP, genetically encoded calcium indicator; HA, histamine; HisCl, Histamine-gated Chloride channel; ICE, Caspase-1/Interleukin-1 converting enzyme; NGM, Nematode Growth Medium; NREM, (non-

29 Rapid Eye Movement); n.s., not significant; R, fluorescence of GCaMP divided by
30 fluorescence of mKate2; RIM, Ring Interneuron M class name; RIS, Ring Interneuron S
31 class name; PVC, Posterior Ventral Cord neuron of the lumbar ganglion class name; R2,
32 ring neurons of the ellipsoid body; ReaChR, Red-activatable channelrhodopsin; SDQL,
33 Posterior lateral interneuron class name - left cell; URY, neuron class name.

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37

38 **Abstract**

39

40 Sleep-active neurons depolarize during sleep to suppress wakefulness circuits. Wake-
41 active wake-promoting neurons in turn shut down sleep-active neurons, thus forming a
42 bipartite flip-flop switch. However, how sleep is switched on is unclear because it is not
43 known how wakefulness is translated into sleep-active neuron depolarization when the
44 system is set to sleep. Using optogenetics in *Caenorhabditis elegans*, we solved the
45 presynaptic circuit for depolarization of the sleep-active RIS neuron during
46 developmentally regulated sleep, also known as lethargus. Surprisingly, we found that RIS
47 activation requires neurons that have known roles in wakefulness and locomotion behavior.
48 The RIM interneurons—which are active during and can induce reverse locomotion—play
49 a complex role and can act as inhibitors of RIS when they are strongly depolarized and as
50 activators of RIS when they are modestly depolarized. The PVC command interneurons,
51 which are known to promote forward locomotion during wakefulness, act as major
52 activators of RIS. The properties of these locomotion neurons are modulated during
53 lethargus. The RIMs become less excitable. The PVCs become resistant to inhibition and
54 have an increased capacity to activate RIS. Separate activation of neither the PVCs nor the
55 RIMs appears to be sufficient for sleep induction; instead, our data suggest that they act in
56 concert to activate RIS. Forward and reverse circuit activity is normally mutually
57 exclusive. Our data suggest that RIS may be activated at the transition between forward
58 and reverse locomotion states, perhaps when both forward (PVC) and reverse (including
59 RIM) circuit activity overlap. While RIS is not strongly activated outside of lethargus,
60 altered activity of the locomotion interneurons during lethargus favors strong RIS
61 activation and thus sleep. The control of sleep-active neurons by locomotion circuits
62 suggests that sleep control may have evolved from locomotion control. The flip-flop sleep
63 switch in *C. elegans* thus requires an additional component, wake-active sleep-promoting
64 neurons that translate wakefulness into the depolarization of a sleep-active neuron when
65 the worm is sleepy. Wake-active sleep-promoting circuits may also be required for sleep
66 state switching in other animals, including in mammals.

67

68 **Introduction**

69

70 Sleep is a behavior that affects many, if not all, physiological processes. Disorders and
71 curtailment of sleep affect the lives of 10% to 30% of the adult population of modern
72 societies. Sleep loss is associated with an increased risk of infection [1], cardiovascular
73 disease [1], psychiatric disease (including depression [2,3]), obesity [4,5], type 2 diabetes
74 [4,5], and cancer [1]. The high prevalence of insomnia and insufficient sleep quality thus
75 presents a massive unmet health and economic problem [1,3-5]. To understand how sleep
76 behavior is generated, it is crucial to solve the underlying neural circuits.

77

78 Sleep circuits require inhibitory sleep-active sleep-promoting neurons, which depolarize
79 specifically at sleep onset and actively induce sleep by releasing inhibitory
80 neurotransmitters, GABA and neuropeptides, to dampen arousal and the activity of wake
81 circuits [6]. Sleep behavior induced by inhibitory sleep-active neurons includes the
82 suppression of voluntary movements and sensory perception, reversibility, and
83 homeostasis [7]. Inhibitory sleep-active neurons suppress wake circuits and can be rapidly
84 suppressed by arousing stimulation to allow for quick awakening. Forced wakefulness is
85 followed by an increase of sleep-active neuron depolarization, which leads to homeostatic
86 sleep corrections. Thus, understanding sleep control requires comprehension of the circuit
87 mechanisms that determine when and how much inhibitory sleep-active neurons depolarize
88 [6,8].

89

90 Circuits control the depolarization of inhibitory sleep-active neurons. For example, wake-
91 active wake-promoting neurons promote arousal and suppress inhibitory sleep-active
92 neurons, whereas sleep need causes sleep-active neuron depolarization. Thus, sleep-active
93 sleep-promoting and wake-active wake-promoting neurons form a flip-flop switch, which
94 ensures that sleep and wake exist as discrete states. This sleep switch is under the control
95 of arousal that favors wake and inhibits sleep through the suppression of sleep-active
96 neurons by inhibitory wake-active neurons [6,9]. It has been proposed that sleep induction
97 is favored by disinhibition of inhibitory sleep-active neurons [10-12]; also, excitatory
98 sleep-active neurons exist that might perhaps present activators of inhibitory sleep-active

99 neurons [13]. However, the forces and mechanisms that flip the sleep switch from wake to
100 sleep when an organism gets sleepy cannot be satisfactorily explained by the present circuit
101 models as it is unclear how sleep-active neurons are turned on when the system is set to
102 sleep.

103

104 Sleep is under circadian and homeostatic controls that determine the timing of sleep and
105 ensure that enough of this essential physiological state takes place [14]. Sleep homeostasis
106 comprises multiple mechanisms that act on different timescales. On long timescales, sleep
107 is a function of prior wakefulness, i.e., prolonged wakefulness leads to increased sleep
108 propensity, and sleep loss triggers compensatory increases in the intensity or duration of
109 sleep. This chronic sleep homeostasis likely is mediated by several parallel mechanisms.
110 For example, in mammals, somnogens such as adenosine accumulate during wakefulness,
111 leading to the inhibition of wake-promoting neurons [15,16]. In *Drosophila*, activity-
112 dependent plasticity of sleep-promoting neurons increases during wakefulness to increase
113 subsequent sleep [17,18]. On short timescales, acute homeostasis determines whether the
114 system's actual state matches the system's set point and carries out corrective action if
115 those values do not match. For example, to homeostatically maintain sleep despite
116 disturbance, micro-arousals need to be compensated for. In humans, homeostatic sleep
117 maintenance can be seen in electroencephalogram (EEG) recordings in the form of k-
118 complexes, in which a spontaneous or evoked short cortical up state is followed by a down
119 state [19-21]. Homeostatic sleep maintenance is also found during sleep in *C. elegans*, in
120 which sleep bouts are interrupted by short motion bouts, with the length of a motion bout
121 correlating with the length of the subsequent sleep bout [22,23]. Thus, across systems,
122 homeostatic sleep maintenance requires constant surveillance of sleep and corrective
123 action.

124

125 Sleep-active sleep-promoting neurons are conserved regulators of sleep and have been
126 found both in vertebrates as well as in invertebrates [8,24]. Mammals possess several
127 populations of sleep-active neurons, most of which are inhibitory, across the brain. These
128 neurons reside in the anterior hypothalamus, brain stem, and cortex [6,12]. Excitatory
129 sleep-active neurons were found in the perioctular midbrain that project to inhibitory sleep-

130 active neurons in the anterior hypothalamus, the role of which could be to activate
131 inhibitory sleep-active neurons [13]. Studying sleep in less complex brains facilitates sleep
132 circuit analysis. In *Drosophila*, sleep-promoting neurons are found at several locations in
133 the brain. A well-characterized population of sleep-promoting neurons is formed by
134 neurons residing in the dorsal fan-shaped body (dFB). R2 ring neurons of the ellipsoid
135 body accumulate homeostatic sleep pressure over time to promote activation of sleep-
136 promoting dFB neurons, probably by an indirect mechanism [17,18]. *C. elegans* possesses
137 a single inhibitory sleep-active neuron called RIS (Ring Interneuron S class name). Like
138 its mammalian counterparts, RIS depolarizes at sleep onset. RIS is crucial for sleep
139 induction because its ablation leads to a virtually complete loss of detectable sleep bouts
140 [25-27]. The small, invariant nervous system, its mapped connectome, and the transparency
141 of *C. elegans* facilitate neural circuit analysis [28]. However, the specific neural circuits
142 that control RIS activity are not yet understood.

143

144 *C. elegans* shows sleep behavior during many stages and conditions. Here, we analyzed
145 sleep behavior during development, also known as lethargus, the stage prior to each of the
146 4 molts during larval development [8,27,29-31]. We used optogenetics to dissect the neural
147 circuits that control the activation of the sleep-active RIS neuron in *C. elegans*. We found
148 a third and novel important element of the flip-flop switch: interneurons that are active
149 during wakefulness and that are known to control locomotion are required for RIS
150 activation and sleep. These findings suggest a tripartite flip-flop circuit model that can
151 explain how arousing stimulation inhibits RIS depolarization, how RIS depolarization is
152 homeostatically controlled, and how reduced arousal can induce RIS depolarization. Our
153 RIS circuit model has 2 important implications for understanding sleep control: (1) it
154 suggests that sleep control has evolved from circuits controlling locomotion; and (2) sleep
155 induction requires an important third element, wake-active sleep-promoting neurons,
156 which translate wakefulness into sleep when the animal is sleepy but awake.

157

158 **Results**

159

160 **Interneurons known to govern locomotion behavior control RIS activity**

161

162 RIS is crucially required for sleep and typically activates during sleep bouts (Fig 1A) [25].
163 However, the presynaptic driver neurons that activate and control this neuron are not
164 known. To identify the circuits controlling RIS activation, we optogenetically tested the
165 role of neurons that are presynaptic to RIS according to the *C. elegans* connectome [28].
166 The neurons called AVJL, CEPDL, URYVL, RIMR, PVCL, and SDQL have been shown
167 to be presynaptic to RIS [28,32]. To find out how these presynaptic neurons control RIS,
168 we activated them with ReaChR (red-activatable channelrhodopsin) and green light and
169 followed RIS calcium activity using GCaMP (a genetically encoded calcium indicator)
170 during and outside of lethargus. We confirmed the expression of ReaChR through a fused
171 fluorescent reporter (mKate2). AVJ, CEPD, URYV, RIM, PVC, and SDQ each are a pair
172 of 2 neurons, of which only one is presynaptic to RIS. Because only promoters that express
173 in both neurons of each pair are available—and because the 2 neurons of each pair are in
174 close proximity—we always manipulated both neurons of the neuronal pair (except for
175 SDQL) [28,32]. Because there were no specific promoters available for the expression in
176 SDQL and PVC, we expressed ReaChR using semi-specific promoters and selectively
177 illuminated only the presynaptic neuron class. We used L1 larvae for most of the
178 optogenetic experiments to dissect the circuit. As SDQL is born postembryonically and
179 likely is not yet functional during the L1 stage, we used L4 larvae to assay its function [33].
180 We compared the effects of optogenetic stimulation outside and during lethargus, defined
181 as the period prior to the molt during which the animals do not feed [34]. Before lethargus,
182 we measured an activation of RIS upon depolarization of AVJ, CEP, and SDQL. During
183 lethargus, the activation of CEP, PVC, and SDQL caused RIS activation (Figs 1B and S1A
184 Fig).

185

186 **Fig 1. Presynaptic neurons control the activity of the sleep-active RIS neuron. (A)**
187 **Sample trace of RIS activity and worm locomotion behavior outside of and during**
188 **lethargus. RIS has no strong calcium transients outside of lethargus but shows strong**
189 **activity transients during lethargus. Upon RIS activation, worms enter sleep bouts. (S1**
190 **Data, Sheet 1A). (B) Presynaptic neurons activate or inhibit RIS outside of and during**
191 **lethargus. For statistical calculations, neural activities before the stimulation period (0–**

192 0.95 min) were compared to activity levels during the stimulation period (1–1.95 min). $*p$
193 < 0.05 , $**p < 0.01$, $***p < 0.001$, Wilcoxon signed rank test. (S1 Data, Sheet 1B). (C)
194 RIS activity decreases upon optogenetic PVC and RIM hyperpolarization. Statistical
195 calculations were performed as described in panel B, but in experiments in which SDQL
196 was stimulated, baseline activity levels were calculated over the time interval from 0.6 to
197 0.95 min. Baseline activity levels were calculated starting from 0.6 min as baseline
198 activity levels were instable before that time point. $*p < 0.05$, $**p < 0.01$, Wilcoxon
199 signed rank test. (S1 Data, Sheet 1C). (D) Circuit model of the RIS presynaptic
200 regulatory network. Activating synaptic input is shown as green arrows, inhibitory
201 synaptic input is shown as red arrows, and unclear synaptic input is shown as black
202 arrow. Gap junctions are indicated as black connections. Neurons that are presynaptic to
203 RIS present mostly activators. PVC is essential for lethargus-specific RIS activation.
204 RIM can inhibit RIS through tyramine and FLP-18 and can activate RIS with glutamate.
205 AVA, Ventral cord interneuron class name; AVJ, neuron class name; CEP, cephalic
206 neuron class name; $\Delta F/F$, change of fluorescence over baseline; FLP-18, FMRF-Like
207 Peptide 18; GCaMP, genetically encoded calcium indicator; n.s., not significant; PVC,
208 Posterior Ventral cord neuron class name; RIM, Ring Interneuron M class name; RIS,
209 Ring Interneuron S class name; SDQL, Posterior lateral interneuron class name - left cell;
210 URY, neuron class name.

211

212

213 All neurons showed consistent effects on RIS depolarization except RIM. RIM is known
214 to play complex roles in controlling behavior and is involved in seemingly opposing
215 behaviors. For example, specific RIM activation can trigger a reversal [35], whereas RIM
216 inhibition has been suggested to be required for reversals through an alternative circuit
217 [36]. We performed optogenetic depolarization experiments of RIM expressing ReaChR
218 using 2 different promoters, the *tdc-1* promoter, which is known to express strongly, and
219 the *gcy-13* promoter, which is known to express at a lower level [37]. Activation of RIM
220 with channelrhodopsin expressed from the *tdc-1* promoter has previously been shown to
221 cause reversals [35], and we observed that activation of RIM using ReaChR expressed from
222 this promoter led to RIS inhibition (Fig 1B, RIM panel). The *tdc-1* promoter expresses

223 strongly in RIM, but also weakly in RIC [38]. To test whether the inhibitory effect of *tdc-*
224 *1* promoter-driven ReaChR expression on RIS was caused by RIC, we also specifically
225 expressed ReaChR in RIC using the *tbh-1* promoter [38]. Specific RIC activation led to
226 RIS activation rather than inhibition (S1B Fig). Therefore, the *tdc-1::ReaChR*-mediated
227 RIS inhibition appears to stem from RIM activation. Activating RIM with the weaker *gcy-*
228 *13* promoter did not cause any net effects on RIS when all trials were averaged (S1C Fig).
229 Visual inspection of the individual trials, however, showed that RIM activation could either
230 inhibit or activate RIS. We therefore sorted single trials for the *gcy-13* experiment into 2
231 classes, in which RIM either activated or inhibited RIS function (S1D Fig). The activation
232 or inhibition of RIS by RIM was indistinguishable during the beginning or end of lethargus
233 (S1E Fig).

234

235 To confirm that RIM can both activate and inhibit RIS, we tested whether activation and
236 inhibition are mediated by different neurotransmitters. We tested the effects of RIM
237 activation on RIS in mutants, which lack transmitters that are known to be expressed in
238 RIM. The RIM neurons are well known to inhibit downstream neurons using tyramine,
239 which requires the *tdc-1* gene [38], and also express neuropeptides (FMRF-Like Peptide
240 18) encoded by the *flp-18* gene [39]. To test whether RIM can inhibit RIS using these
241 known transmitters, we analyzed mutant worms that lack functional *flp-18* and *tdc-1*.
242 Individual inactivation of *flp-18* and *tdc-1* reduced—and double mutation abolished—the
243 inhibition of RIS by RIM (S2 Fig). Therefore, the transmitters tyramine and FLP-18 are
244 together responsible for RIS inhibition by RIM. We next tested activation of RIS by RIM
245 in *eat-4(ky5)* mutant larvae, which lack glutamatergic signaling in many neurons, including
246 RIM [40,41]. RIS activation by RIM activation was completely gone in *eat-4(ky5)* mutant
247 larvae (S3 Fig, we used L4 larvae for this assay as the response was more robust).
248 Therefore, glutamate is required for RIS activation by RIM. Together, these results suggest
249 that RIM can act both as an activator as well as an inhibitor of RIS by employing different
250 neurotransmitters, with weaker activation allowing for RIS activation and stronger
251 activation favoring inhibition.

252

253 The majority of synaptic inputs into RIS that we studied had activating effects; the sole
254 inhibitory effect was observed after strong activation of RIM, whereas weaker RIM
255 activation could also lead to RIS activation. The CEP, URY, and SDQL neurons present
256 sensory receptors and might play a role in activating RIS in response to stimulation. For
257 example, CEP might activate RIS as part of the basal slowing response [42,43]. The PVCs
258 appeared to be strong activators of RIS specifically during lethargus. This suggests either
259 that the PVC-to-RIS connection might be specific to lethargus or that it has not yet matured
260 during the mid-L1 stage. We therefore repeated the experiment and activated PVC in L2
261 larvae. PVC activated RIS both during and outside of lethargus, but the activation during
262 lethargus was much stronger, suggesting that the activation of RIS by PVC is strongly
263 enhanced during lethargus (S4 Fig).

264

265 To find out which presynaptic neurons are required for inhibition or activation of RIS
266 during lethargus, we tested the effect of optogenetic inhibition of the presynaptic neurons
267 on RIS activation. We used ArchT (archaerhodopsin from *Halorubrum* strain TP009),
268 which hyperpolarizes neurons by pumping protons out of the cell [44,45]. As earlier, we
269 verified the expression of ArchT in neurons of interest by using an mKate2-tagged version.
270 As in the ReaChR experiments, we specifically illuminated each presynaptic neuron class
271 and quantified RIS activation using calcium imaging. Before lethargus, inhibition of AVJ
272 and PVC led to an inhibition of RIS, whereas inhibition of the other neurons tested had no
273 acute statistically significant effect on RIS (optogenetic RIM hyperpolarization using the
274 stronger *tdc-1* promoter in worms outside of lethargus showed a tendency to inhibit RIS
275 function [$p = 0.0539$; $N = 11$ animals], whereas the weaker *gcy-13* promoter had no
276 detectable effect). During lethargus, optogenetic inhibition of PVC and RIM (using the
277 stronger *tdc-1* promoter) led to significant RIS inhibition, whereas there was no effect seen
278 for the other neurons (Fig 1C and S5A Fig; inhibition of RIM using the weaker *gcy-13*
279 promoter only produced a tendency but no statistically significant net effect, S5B and S5C
280 Figs).

281

282 Absence of an effect of optogenetic inhibition of presynaptic neurons could mean either
283 that these neurons are not required for RIS activation, that the inhibition was not strong

284 enough, or that they may act redundantly (we did not find any evidence for redundancy, at
285 least for CEP and URY, S5D Fig). Our optogenetic analysis revealed a complex set of
286 presynaptic inputs for regulation of RIS activity (Fig 1D). The optogenetic depolarization
287 experiments suggest that CEP, PVC, RIM, and SDQL present the most potent presynaptic
288 activators of RIS. The capacity of PVC to activate RIS is strongly increased during
289 lethargus, indicating that this neuron is involved in the lethargus-specific activation of RIS.
290 The optogenetic hyperpolarization experiments suggest that PVC and RIM are essential
291 presynaptic activators of RIS during lethargus. Therefore, we focused our analysis on PVC
292 and RIM neurons.

293

294 **PVC becomes resistant to inhibition during lethargus**

295

296 Neuronal activation and silencing experiments revealed PVC as a main activator of RIS.
297 These results predict that neuronal activity of PVC should correlate with RIS activation
298 and sleep bouts. To test for such correlation, we measured the activity of both neurons
299 simultaneously. Because the calcium transients observable in PVC are typically small [46]
300 and could not be detected in our assays in mobile worms (data not shown), we immobilized
301 the larvae and used RIS activation as a proxy for sleep bouts. We extracted both RIS and
302 PVC activity and aligned all data to the RIS activation maxima. This analysis showed that
303 PVC activated approximately 1 min earlier than RIS and reached its maximum activation
304 approximately 1.5 min earlier than RIS. PVC activity decreased slowly during the RIS
305 transient (Fig 2A). This result is consistent with a role for PVC in promoting RIS
306 depolarization.

307

308 **Fig 2. PVC is an RIS activator that becomes resistant to inhibition during lethargus,**
309 **but PVC activation is not sufficient for sleep induction.** (A) Simultaneous PVC and
310 RIS GCaMP traces aligned to RIS peaks in fixed L1 lethargus worms. PVC activates
311 before the RIS peak and stays active until the peak. $*p < 0.05$, $**p < 0.01$, Wilcoxon
312 signed rank test. (S1 Data, Sheet 2A). (B) PVC hyperpolarization inactivates RIS and
313 induces behavioral activity. PVC hyperpolarization was performed by expressing ArchT
314 under the *zk637.11* promoter. In contrast to the *nmr-1* promoter, the *zk637.11* promoter

315 lacks expression in head command interneurons. $**p < 0.01$, $***p < 0.001$, Wilcoxon
316 signed rank test for GCaMP and speed, Fisher's exact test for sleep fraction. (S1 Data,
317 Sheet 2B). (C) During lethargus, PVC becomes resistant to inhibition. Outside of
318 lethargus, its inhibition is stronger and continues beyond the end of optogenetic
319 stimulation. During lethargus, PVC activity levels return back to baseline already during
320 the stimulation period. $*p < 0.05$, $**p < 0.01$, Wilcoxon signed rank test. (S1 Data, Sheet
321 2C). (D) PVC activation translates into mostly a forward mobilization in L1 lethargus. $*p$
322 < 0.05 , $***p < 0.001$, Wilcoxon signed rank test for Speed. Fisher's exact test for fraction
323 of direction. (S1 Data, Sheet 2D). ArchT, archaerhodopsin from *Halorubrum* strain
324 TP009; ATR, all-trans-retinal; $\Delta F/F$, change of fluorescence over baseline; GCaMP,
325 genetically encoded calcium indicator; n.s., not significant; PVC, Posterior Ventral cord
326 neuron class name; RIS, Ring Interneuron S class name;

327

328 PVC inhibition reduced RIS activity in immobilized animals, but it is unclear how PVC
329 inhibition affects behavior. To be able to test the effects of PVC inhibition on behavior
330 without affecting the other command interneurons, we chose a more specific promoter for
331 expression in PVC from single-cell RNA sequencing data. There was no gene in the
332 available datasets that was expressed only in the cluster of cells containing PVC, but the
333 previously uncharacterized gene *zk673.11* was expressed specifically in PVC and in only
334 a few other neurons excluding other command interneurons [47,48] (personal
335 communication from J. Packard to H. Bringmann. S6 Fig). Hyperpolarization of PVC using
336 ArchT driven by the *zk673.11* promoter led to an acute inhibition of RIS, an increase in
337 locomotion, and a reduction of sleep (Fig 2B). Hyperpolarization of PVC using ArchT also
338 strongly inhibited RIS outside of lethargus. This experiment confirmed the role of the
339 PVCs in activating RIS.

340

341 Hyperpolarization of PVC outside of lethargus appeared to have a stronger and longer-
342 lasting effect on RIS inhibition compared with during lethargus (Figs 1C and 2B). This is
343 surprising because PVC is a stronger activator of RIS during lethargus in the optogenetic
344 activation experiments (Fig 1B). This effect could be explained if PVC responded more
345 severely to inhibition outside of lethargus. We tested this idea by inhibiting PVC using

346 ArchT and green light and simultaneously imaged PVC activity. PVC hyperpolarization
347 was stronger in worms outside of lethargus, and PVC remained inhibited after the
348 optogenetic manipulation. During lethargus, PVC was only weakly inhibited at the
349 beginning of optogenetic stimulation and returned to baseline levels already during the
350 stimulation (Fig 2C). We also tested whether optogenetic excitability of PVC was
351 modulated during lethargus but could not find any differences in excitability of PVC during
352 or outside of lethargus (S7A Fig). Thus, PVC is more susceptible to inhibition outside of
353 lethargus but becomes resistant to inhibition during lethargus. This effect can explain the
354 stronger hyperpolarization of RIS during PVC inhibition outside of lethargus, and this
355 effect likely presents an important modulation of the circuit to favor PVC activation and
356 thus RIS activation during lethargus.

357

358 PVC is known to promote forward movement upon posterior mechanical stimulation, and
359 optogenetic stimulation of PVC in adults has been shown to promote forward locomotion
360 [49,50]. Our data showed that PVC also activates the RIS neuron, and consistent with this
361 observation, mechanical stimulation caused RIS activation (S7B Fig). This suggests that
362 PVC activates RIS to modulate forward locomotion speed and to promote sleep. However,
363 it is unclear how PVC can promote forward motion and sleep, as these are two seemingly
364 opposing behaviors. We therefore tested whether optogenetic stimulation of PVC in larvae
365 induces sleep behavior. We activated PVC using *nmr-1::ReaChR* in mobile L1 larvae
366 during lethargus and specifically illuminated the tail of the animal, which contains the cell
367 bodies of the PVC neurons but not the other *nmr-1*-expressing neurons. We quantified the
368 speed as well as the direction of movement of the worm. During PVC activation during
369 lethargus, the worms visibly accelerated movement and mostly crawled forward, but we
370 could not see induction of sleep behavior during optogenetic stimulation (Fig 2D).
371 Consistent with this finding, optogenetic PVC activation during and before lethargus
372 always led to the activation of AVB (Ventral cord interneuron class name) interneurons,
373 which are known to be premotor neurons required for forward locomotion [49] (S7C and
374 S7D Figs). Together, these experiments showed that PVC activates prior to RIS and is
375 required for RIS activation. However, its activation alone does not seem to be sufficient to
376 induce sleep behavior.

377

378 **RIS and PVC activate each other forming a positive feedback loop**

379

380 PVC presents a major activator of RIS, but how a forward command interneuron can cause
381 strong and state-specific activation of the RIS neuron during sleep bouts is not clear. We
382 therefore tested how optogenetic RIS activation affects PVC activity. We selectively
383 activated RIS using ReaChR and measured calcium activity in PVC in immobilized
384 animals. Upon RIS stimulation, PVC immediately displayed unexpectedly strong calcium
385 transients, which were slightly stronger during lethargus (Fig 3A and S8A Fig). These
386 results show that PVC and RIS activate each other, thus forming a positive feedback loop.
387 The sleep-inducing RIS neuron has so far only been shown to inhibit other neurons, making
388 PVC the first neuron that is not inhibited but is activated by RIS. For example, command
389 interneurons such as AVE and AVA (ventral cord interneuron class names) and other
390 neurons are not activated but are inhibited by RIS [25].

391

392 **Fig 3. RIS and PVC activate each other, forming a positive feedback loop.** (A–E) RIS
393 depolarization leads to a strong PVC depolarization outside of and during lethargus. This
394 PVC depolarization is almost abolished in *flp-11(tm2705)*, and it is significantly reduced
395 in AVE-ablated worms. $*p < 0.05$, $**p < 0.01$, Wilcoxon signed rank test (S1 Data,
396 Sheets 3A, 3B, 3C-E). (F) AVE-ablated worms show increased sleep. AVA-ablated
397 worms do not show a significant sleep phenotype. Shown are sleep fractions during
398 lethargus. $*p < 0.05$, Kolmogorov-Smirnov test (S1 Data, Sheet 3F). (G) RIS does not
399 reach the same activation levels in *aptf-1(gk794)* and *flp-11(tm2705)* mutants compared
400 to wild-type worms. *aptf-1(gk794)* and *flp-11(tm2705)* mutants neither immobilize nor
401 sleep during RIS activation. $***p < 0.001$, Welch test (S1 Data, Sheet 3G-I). (H) *flp-*
402 *11(tm2705)* mutants have significantly fewer wide RIS peaks. *aptf-1(gk794)* mutants
403 display the same amount of wide RIS peaks as wild-type worms. $**p < 0.01$,
404 Kolmogorov-Smirnov test (S1 Data, Sheet 3G-I). (I) *flp-11(tm2705)* and *aptf-1(gk794)*
405 mutants do not show sleep during lethargus. $**p < 0.01$, Kolmogorov-Smirnov test (S1
406 Data, Sheet 3G-I). (J) A circuit model for the positive feedback loop between RIS and
407 PVC. Activating synaptic input is shown as green arrows, inhibitory synaptic input is

408 shown as red arrows, and gap junctions are indicated as black connections. During
409 wakefulness, reverse command interneurons inhibit PVC so that PVC does not activate
410 RIS. During lethargus, PVC directly activates RIS, which then inhibits reverse command
411 interneurons through FLP-11. This may speculatively disinhibit PVC, leading to a
412 positive feedback. AVA, AVE, PVC, RIS, neuronal class names; $\Delta F/F$, change of
413 fluorescence over baseline; FLP-11, FMRF-Like Peptide 11; GCaMP, genetically
414 encoded calcium indicator; n.s., not significant;

415

416 RIS induces sleep through the release of neuropeptides with the major sleep-inducing
417 neuropeptides encoded by the *flp-11* gene [51]. To test whether FLP-11 neuropeptides are
418 required for RIS-induced PVC activation, we repeated the optogenetic RIS activation with
419 simultaneous PVC calcium measurement in an *flp-11* deletion mutant. RIS-induced PVC
420 activation was almost completely abolished in the *flp-11* deletion (reduction of transient
421 maximum by 79% during lethargus), indicating that FLP-11 neuropeptides are required for
422 RIS-induced PVC activation (Fig 3B).

423

424 While PVC is presynaptic to RIS, RIS is not presynaptic to PVC [28,32]. The activation of
425 PVC by RIS could involve diffusional mechanisms or could be indirect through other
426 neurons, perhaps mediated by the inhibition of a PVC inhibitor such as AVA/AVD/AVE.
427 RIS has been shown to inhibit AVA/AVE [25], and RIS is presynaptic to AVE [28,32],
428 suggesting that PVC activation involves inhibition of AVE. We therefore repeated RIS
429 activation and PVC calcium imaging in a strain in which AVE was impaired through
430 expression of tetanus toxin [52]. The initial PVC activation maximum after AVE
431 impairment was reduced by 43% during lethargus, but subsequent PVC activity was
432 increased (Fig 3C-E). AVE is connected to other reverse command interneurons, which
433 collectively inhibit PVC [28,53]. This circuit design suggests that AVE might play a dual
434 role in controlling RIS activity. It should have a positive role in mediating activation of
435 PVC through RIS and thus could promote the feedback loop, but it should also have an
436 inhibiting role by promoting PVC inhibition. To test for a role of the arousal neurons AVE
437 and AVA in sleep, we inhibited AVE with tetanus toxin [52] and AVA using HisCl
438 (Histamine-gated Chloride channel) [54] and quantified sleep amount. Whereas we could

439 not find any effect of AVA impairment on sleep amount, AVE impairment led to an
440 average increase of sleep by 42% (Fig 3F). Together, these data suggest that PVC and RIS
441 rely on positive feedback for their activation that involves the release of FLP-11
442 neuropeptides and inhibition of PVC by AVE.

443

444 If depolarization of RIS activates PVC, what consequences does hyperpolarization of RIS
445 have on PVC activity? To answer this question, we measured the response of PVC to RIS
446 inhibition. We hyperpolarized RIS optogenetically for 1 min using ArchT and measured
447 the activity of PVC. Interestingly, PVC showed a small but significant activity increase
448 during RIS inhibition, an effect that was increased during lethargus (S8B Fig). The
449 disinhibition of PVC by RIS inactivation is likely not direct and may reflect a general
450 increase in neuronal and behavioral activity that is caused by RIS inhibition and that
451 extends to the PVC neurons. Because PVC is a major activator of RIS, its disinhibition
452 could be part of a homeostatic feedback regulation.

453

454 Our results suggest that there is a positive feedback from sleep induction onto RIS
455 activation and that full RIS activation is only possible when sleep is successfully induced,
456 explaining the strong correlation of RIS depolarization and sleep-bout induction [27]. This
457 model would predict that RIS transients are dampened if RIS is not able to induce sleep
458 bouts. To test this idea, we analyzed RIS calcium transients in *aptf-1(-)* mutant worms in
459 which RIS still shows depolarization transients during lethargus but cannot efficiently
460 induce quiescence [25,51]. In *aptf-1(-)* mutant animals, calcium transient maxima were
461 reduced by about 35% (Fig 3G-I). A major function of APTF-1 (Activating enhancer
462 binding Protein 2 Transcription Factor 1) is the expression of FLP-11 neuropeptides that
463 are required for quiescence induction [51]. To test whether FLP-11 neuropeptides play an
464 essential role in shaping RIS transients, we measured RIS calcium transients in mutant
465 worms carrying a deletion of *flp-11*. These mutant animals showed only a reduced number
466 of long RIS transients that were of reduced size (Fig 3G-I). *flp-11(-)* showed, however,
467 many short RIS transients (S8C-F Fig) that were not associated with sleep bouts but may
468 reflect attempts to induce sleep bouts. These results are consistent with the idea that sleep

469 induction is a self-enforcing process in which RIS-mediated inhibition of brain activity
470 through FLP-11 neuropeptides promotes long RIS calcium transients (Fig 3J).

471

472 We next tested what feedback interaction exists between RIM and RIS neurons. We
473 optogenetically depolarized or hyperpolarized RIS and measured RIM activity. RIS
474 activation did not significantly change RIM activity, but there was a small inhibitory trend
475 (S8G Fig). RIS inhibition led to an activation of RIM (S8H Fig). These results show that,
476 while RIM can activate as well as inhibit RIS, RIS is an inhibitor of RIM.

477

478 **RIM can activate RIS, but its activation is not sufficient for sleep induction**

479

480 A second important activator of RIS is RIM. We therefore asked whether RIM, similar to
481 PVC, also is active prior to RIS depolarization and sleep bouts. We measured RIM activity
482 by imaging GCaMP in moving worms. All sleep bouts were extracted, and RIM activity
483 was aligned to sleep-bout onset. Averaged RIM activity peaked approximately 30 s before
484 the beginning of the sleep bout (Fig 4A). This finding is consistent with a function for RIM
485 in RIS activation. We then asked whether RIM is required for sleep induction. We ablated
486 RIM through expression of *egl-1* under the *tdc-1* promoter. We quantified lethargus sleep
487 in RIM-ablated worms. RIM-ablated larvae showed a normal fraction of sleep, a slightly
488 increased frequency of sleep bouts, and a normal length of sleep bouts (Fig 4B-D). In
489 analogy to the PVC experiments, we analyzed the effect of optogenetic RIM depolarization
490 on behavior. We first tested behavior caused by activation of RIM with ReaChR driven by
491 the strong *tdc-1* promoter on the locomotion of worms. Consistent with previous findings
492 [35] and our observation that RIS is inhibited under these conditions, RIM activation
493 during lethargus caused mobilization, and larvae crawled mostly backwards (Fig 4E). We
494 next tested for the effects of weaker RIM activation using the *gcy-13* promoter. Activation
495 of RIM caused increased mobility when RIS was inhibited. In trials in which RIM
496 activation led to RIS activation, there was no significant change of speed of the worms
497 (S1D Fig). We next wanted to test whether excitability of RIM is altered during the
498 lethargus state. We therefore activated RIM strongly using the *tdc-1* promoter and
499 measured RIM activity. Outside of lethargus, RIM was strongly excited. During lethargus,

500 however, excitability was strongly reduced (Fig 4F and 4G). In summary, RIM activation
501 is not sufficient to induce sleep. RIM could, however, contribute to strong RIS activation
502 and sleep induction by acting in concert with other neurons. Reduced excitability of RIM
503 during lethargus could favor the activating effect of RIM on RIS while dampening the
504 inhibiting effects of RIM on RIS.

505

506 **Fig 4. RIM activity peaks prior to sleep bouts, but RIM activation is not sufficient**
507 **for sleep induction.** (A) RIM activates prior to sleep bouts. $*p < 0.05$, Wilcoxon signed
508 rank test (S1 Data, Sheet 4A). (B-D) RIM-ablated worms have an increased sleep-bout
509 frequency, while the sleep fraction and bout duration are not significantly changed during
510 L1 lethargus. RIM was genetically ablated by expressing *egl-1* under the *tdc-1* promoter.
511 $*p < 0.05$, Kolmogorov-Smirnov test (S1 Data, Sheet 4B-D). (E) RIM depolarization
512 leads to increased mobility and reverse motion. $*p < 0.05$, $***p < 0.001$, Wilcoxon
513 signed rank test for speed. Fisher's exact test for fraction of direction (S1 Data, Sheet
514 4E). (F-G) During lethargus, RIM becomes resistant to activation. RIM was
515 optogenetically activated using ReaChR expressed under the *tdc-1* promoter. Outside of
516 lethargus, its activation is stronger (F). Activity levels during the stimulation period were
517 quantified by subtracting baseline activity levels from levels during the stimulation
518 period (G). $*p < 0.05$, $**p < 0.01$, Wilcoxon signed rank test for GCaMP and
519 Kolmogorov-Smirnov test for quantification of stimulation levels (S1 Data, Sheet 4F-G).
520 ATR, all-trans-retinal; $\Delta F/F$, change of fluorescence over baseline; GCaMP, genetically
521 encoded calcium indicator; n.s., not significant; ReaChR, red-activatable
522 channelrhodopsin; RIM, Ring Interneuron M class name.

523

524

525

526 **Interneurons regulating locomotion act in concert to activate RIS**

527

528 Separate activation of PVC or RIM neurons caused moderate RIS activation but not the
529 strong activation of RIS that is typically associated with sleep bouts. Thus, hypothetically,
530 multiple neurons act in concert to cause strong RIS activation. Our earlier presynaptic

531 neuron analysis suggests that this hypothetical set of neurons should include PVC and RIM
532 interneurons but could also include additional neurons. Our analysis of RIM and PVC
533 points to neurons of the command interneuron circuit for RIS activation, and thus we tested
534 the effects of ablation of a large fraction of the interneurons controlling locomotion. The
535 *nmr-1* promoter expresses in AVA, AVE, AVD, and PVC command interneurons as well
536 as in second-layer RIM neurons [55]. We used a strain that ablates these locomotion-
537 controlling interneurons by expressing the pro-apoptosis regulator ICE (Caspase-
538 1/Interleukin-1 converting enzyme) from the *nmr-1* promoter [55] and measured sleep and
539 RIS activation. Command interneuron ablation reduced sleep bouts during lethargus by
540 about 76% (Fig 5A), and RIS activation was reduced by 63% (S9A Fig). The movement
541 of command interneuron-ablated worms also was slower (S9B Fig). Quiescence bouts did
542 not occur at the beginning of the lethargus phase as defined by cessation of feeding and
543 were only observed around the middle of the lethargus phase (S9C Fig). An independently
544 generated strain that ablates command interneurons using *egl-1* expression—also by using
545 the *nmr-1* promoter—caused a reduction of sleep by 81% (Fig 5A).

546

547 **Fig 5. The locomotion interneuron circuit controls RIS activation and sleep. (A)**
548 **Command interneurons are responsible for the majority of sleep. Command interneurons**
549 **were genetically ablated by expressing ICE or *egl-1* under the *nmr-1* promoter.**
550 **Command interneurons-ablated worms display a massive loss-of-sleep phenotype. *** p**
551 **< 0.001, Welch test (S1 Data, Sheet 5A). (B) Hyperpolarization of command interneurons**
552 **causes RIS inhibition and suppresses sleep. During lethargus, the hyperpolarization is**
553 **followed by a strong post-stimulation activation of RIS. ** p < 0.01, *** p < 0.001,**
554 **Wilcoxon signed rank test for GCaMP and speed, Fisher's exact test for sleep fraction**
555 **(S1 Data, Sheet 5B). $\Delta F/F$, change of fluorescence over baseline; GCaMP, genetically**
556 **encoded calcium indicator; ICE, Caspase-1/Interleukin-1 converting enzyme; n.s., not**
557 **significant; RIS, Ring interneuron S class name;**

558

559

560 Next, we wanted to test conditional loss of function of the command interneuron circuit on
561 RIS activity. We expressed ArchT broadly in locomotion-controlling interneurons by using

562 the *nmr-1* promoter. We then inhibited all command interneurons using green light and
563 simultaneously imaged the activity of RIS. Inhibition of *nmr-1*-expressing neurons
564 strongly inhibited RIS inhibition both outside and during lethargus. Interestingly, there was
565 a strong post-stimulus activation of RIS, which was strongly increased only during
566 lethargus. This activation peaked at approximately 170% of the RIS baseline. Sleep was
567 inhibited by command interneuron inhibition, and worms reached mobility speeds similar
568 to those outside of lethargus (Fig 5B). Mosaic analysis of an extrachromosomal array
569 carrying the *nmr-1::ArchT* transgene revealed that RIS was partially inhibited when ArchT
570 was expressed in head neurons but not in PVC and that the effect of inhibition was
571 substantially stronger when ArchT was not only expressed in head neurons but also
572 expressed in PVC (S9D-E Fig). This experiment showed that multiple interneurons act in
573 concert to activate RIS and induce sleep. Among the *nmr-1*-expressing interneurons, only
574 RIM and PVC are presynaptic to RIS [28,32]. However, additional reverse command
575 interneurons could also contribute to RIS regulation through indirect mechanisms.

576

577 Because the command interneuron circuit is controlled by glutamatergic signaling [55,56]
578 and because RIM activation of RIS requires glutamate (S3 Fig), we also analyzed the sleep
579 behavior of *eat-4(ky5)* mutant larvae that have impaired glutamatergic neurotransmission.
580 In *eat-4(ky5)* mutant larvae, sleep-bout duration was significantly reduced, whereas sleep
581 bouts occurred with normal frequency. This indicates that glutamate signaling might play
582 a role in the maintenance but not in the initiation of sleep bouts (S10A-D Fig). Consistent
583 with these findings, glutamate signaling also plays a role in the maintenance of NREM
584 (non-Rapid Eye Movement) sleep in mice [13]. *nmr-1(ak4)* glutamate receptor mutant
585 larvae only displayed slightly reduced RIS activation transients, which indicates that
586 additional glutamate receptors are required for sleep induction (S10E-I Fig). Together,
587 these mutant phenotypes support the view that excitatory neurotransmitter systems that are
588 associated with locomotion are important for RIS activation.

589

590 **RIS inhibition causes homeostatic rebound activation**

591

592 The design of the sleep circuit suggests an intimate mutual control mechanism of RIS and
593 command interneurons that could allow homeostatic control of sleep. Arousing stimulation
594 is known to inhibit sleep-active neurons and to increase subsequent sleep [22,23,25,27].
595 Consistent with these published data, we observed that the maximum RIS GCaMP intensity
596 increased logistically with the length of the preceding motion bout during lethargus (S11A
597 Fig). We thus hypothesized that stimulation inhibits RIS and leads to its subsequent
598 depolarization, forming a homeostat that allows maintaining or reinstating sleep bouts. We
599 tested this hypothesis by arousing the worms with a blue light stimulus (Fig 6A-B). During
600 the stimulus, worms mobilized, and sleep was inhibited. In some of the trials, worms went
601 back to sleep promptly after the stimulation and decreased their motion speed again within
602 3 min. Because worms did not remain mobile after the stimulation, we classified these trials
603 as “nonmobilizing.” In these nonmobilizing trials, RIS showed a post-stimulus activation,
604 which was 34% stronger than the baseline activity. RIS activation correlated with a
605 significantly increased fraction of sleep. In other trials during lethargus, the worms stayed
606 mobile for at least 3 min after stimulation and did not go back to sleep. Because worms
607 remained mobile after the stimulation, we classified these trials as “mobilizing.” In these
608 mobilizing trials, RIS stayed inhibited and was 16% less active than the baseline before
609 stimulation (Fig 6A). To measure global neuronal activity during the blue-light stimulation
610 experiment, we imaged worms that expressed pan-neuronal GCaMP [57]. Trials were again
611 divided into mobilizing and nonmobilizing trials during lethargus depending on the
612 mobilization status after the stimulus. Nonmobilizing trials showed a global neuronal
613 inhibition that was 93% of the baseline activity (Fig 6B). These experiments show that
614 noxious blue-light stimulation inhibits sleep and RIS and causes a reactivation of RIS when
615 the system returns to sleep.

616

617 **Fig 6. RIS inhibition causes homeostatic rebound activation.** (A–B) A blue light
618 stimulus leads to awakening and mobilization of *C. elegans*. Worms that go back to sleep
619 after the stimulus show an activation rebound: pan-neuronal inhibition below baseline
620 levels and RIS activation above baseline levels; “lethargus mobilizing” refers to animals
621 that stayed awake and active during the post-stimulus time; “lethargus nonmobilizing”
622 refers to animals that went back to sleep after the stimulation. * $p < 0.05$, ** $p < 0.01$,

623 *** $p < 0.001$, Wilcoxon signed rank test for GCaMP and speed, Fisher's exact test for
624 sleep fraction (S1 Data, Sheet 6A and 6B). (C) RIS shows rebound activation following
625 hyperpolarization. Behavioral and brain activity measurements correlate throughout the
626 whole experiment. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, Wilcoxon signed rank test for
627 GCaMP and speed, Fisher's exact test for sleep fraction (S1 Data, Sheet 6C). (D–E)
628 Dose-response curve of optogenetic RIS hyperpolarization with different stimulus
629 lengths. RIS activation rebound transients saturate with increasing length of inhibition.
630 Worms not showing a rebound activation transient after RIS optogenetic
631 hyperpolarization were excluded from the analysis. Numbers of worms not responding
632 were as follows: (1) In experiments in which RIS was optogenetically inhibited for 48 s,
633 all worms showed an RIS rebound activation transient. (2) In experiments in which RIS
634 was optogenetically inhibited for 5 min, 1 out of 7 worms did not show a RIS rebound
635 activation transient. (3) In experiments in which RIS was optogenetically inhibited for 10
636 min, 1 out of 13 worms did not show an RIS rebound activation transient. Curve in D was
637 fitted as an asymptotic function, and curve in E was fitted as a BoxLucas1 function (S1
638 Data, Sheet 6D, E). $\Delta F/F$, fluorescence change over baseline; GCaMP, genetically
639 encoded calcium sensor; n.s., not significant; RIS, Ring Interneuron S class name; R,
640 fluorescence of GCaMP / fluorescence of mKate2;;

641

642 In normal sleep and in the sensory stimulation experiment, periods of inactivity of RIS
643 were always followed by periods of RIS activation. This suggested that inhibition of RIS
644 causes its subsequent reactivation. We tested this hypothesis by optogenetically
645 hyperpolarizing RIS and following its activity using calcium imaging. We inhibited RIS
646 directly for 60 s by expressing the light-driven proton pump ArchT specifically in this
647 neuron and used green light illumination to activate ArchT. We followed RIS calcium
648 activity using GCaMP during the experiment and quantified behavior. Optogenetic
649 hyperpolarization of RIS led to a decrease in intracellular calcium and increased behavioral
650 activity. Approximately 1 min after the end of the inhibition, RIS showed a rebound
651 activation transient during which calcium activity levels increased strongly and rose well
652 above baseline levels, concomitant with a decrease in behavioral activity. Overall brain
653 activity measurements showed that behavioral activity and brain activity correlated

654 throughout the experiment (Fig 6C). Rebound activation was observed neither following
655 PVC nor following RIM inhibition (Fig 2C and S11B Fig), suggesting that rebound
656 activation is specific to RIS and is not a general property of all neurons [58]. Strikingly,
657 while the rebound transient was also measurable outside of lethargus, the strength of the
658 RIS rebound depolarization was 3-fold stronger during lethargus than before lethargus,
659 indicating that the propensity for RIS rebound activation is strongly increased during
660 lethargus.

661

662 To test whether rebound activation of RIS mediates acute or chronic homeostasis, we tested
663 whether the strength of the rebound activation is a function of length of prior inhibition.
664 For this experiment, we increased the length of the RIS inhibition and quantified the time
665 it took after the end of the stimulation until the rebound transient started as well as the peak
666 maximum of the rebound. After inhibiting RIS for 5 min, the rebound initiated immediately
667 after the end of the stimulation and the maximum that was reached exceeded that observed
668 after about 1 min of RIS stimulation. Inhibiting RIS for 10 min did not further increase the
669 occurrence or strength of the rebound transient. These results show that RIS activation
670 rebound transients rapidly saturate with increasing length of inhibition (Fig 6D-E and
671 S11C-E Fig). Thus, RIS shows a rebound activation following inhibition. The rebound
672 activation presents the translation of RIS inhibition into subsequently increased RIS
673 activity and thus sleep induction. Rebound activation of RIS does not seem to constitute a
674 chronic integrator of wake time but presents an acute homeostatic regulatory phenomenon
675 to induce or reinstate sleep bouts.

676

677 Rebound activation of RIS could present a cell-intrinsic property or could be generated by
678 a neural circuit. To discriminate between these hypotheses, we measured rebound
679 activation in *unc-13(s69)* mutant animals in which synaptic signaling is globally impaired
680 [59], or in worms that express tetanus toxin [60] specifically in RIS to abrogate synaptic
681 transmission specifically in this neuron. Rebound activation of RIS was abolished in
682 RIS::tetanus toxin (S11F-G Fig) as well as *unc-13(s69)* worms (S11F and S11H Fig).
683 These results indicate that rebound activation of RIS is a property of the neuronal network.

684

685 In analogy to the activation rebound seen after optogenetic RIS inhibition, optogenetic RIS
686 activation might cause a negative rebound, i.e., an inhibition of RIS inhibition below
687 baseline levels following its optogenetic activation. Indeed, we observed such an effect.
688 Interestingly, the negative rebound was 3-fold stronger during lethargus compared to
689 outside of lethargus (S11I Fig). However, such a negative rebound was also present in other
690 neurons such as PVC (S7A Fig), making it difficult to judge whether this effect is part of
691 a specific sleep homeostatic system or rather a general response of neurons to strong
692 depolarization [58]. In summary, RIS activity is homeostatically regulated, with its
693 inhibition causing its reactivation. This rebound activation is strongly increased during
694 lethargus and likely is required for inducing or reinstating sleep.

695

696 **Modest dampening of brain arousal occurs upstream of RIS**

697

698 Our results demonstrate that the command interneuron circuit, including PVC, plays a
699 major role in activating RIS involving self-enforcing positive feedback, resulting in strong
700 RIS activation and thus sleep induction. RIS calcium transients are small during
701 development outside of lethargus, whereas transients are high during lethargus. What
702 determines that RIS calcium transients are limited outside of lethargus but promoted during
703 lethargus? As an important principle of command interneuron control, forward and reverse
704 command interneurons inhibit each other to allow discrete forward and reverse locomotion
705 states. The AVA/AVD/AVE/RIM interneurons initiate reverse locomotion by activating
706 premotor interneurons while inhibiting the forward command circuit including AVB/PVC.
707 By contrast, during forward movement, reverse command interneurons are inhibited
708 [49,56].

709

710 Small changes in arousal and activity of the command interneurons can change the
711 equilibrium of forward and reverse command interneurons [55]. Hyperactive mutants
712 suppress sleep across species, including *C. elegans* [61-68]. Many arousal cues trigger
713 backwards escape movements and inhibit RIS [25,27,69]. Thus, previous studies on the
714 command interneuron circuit together with our results suggest that arousal inhibits RIS
715 through inhibiting PVC. This model of RIS activation would predict that there are changes

716 during lethargus that are upstream of RIS activity that change the properties of the
717 command circuit, leading to increased PVC and thus RIS activation.

718

719 We reasoned that it should be possible to measure these changes that occur in command
720 interneuron activity upstream of RIS by characterizing neural activity and behavior in *aptf-*
721 *I(-)* mutant worms. We quantified behavior and command interneuron calcium levels
722 across lethargus in *aptf-1(-)* mutant worms. Wild-type animals showed successive sleep
723 bouts and a 72% reduction in locomotion speed during lethargus. By contrast, *aptf-1(-)*
724 mutant animals almost never showed quiescence bouts (Fig 3I), but nevertheless,
725 locomotion speed was decreased by 20% during the lethargus phase (Fig 7). Consistent
726 with the behavioral activity reduction, there was a significant reduction of command
727 interneuron activity during lethargus also in *aptf-1(-)* mutant animals (Fig 7 and S12 Fig).
728 To further characterize the neuronal changes upstream of RIS-mediated sleep induction,
729 we imaged the activity of RIM during lethargus in *aptf-1(-)* mutants. In wild-type animals,
730 RIM regularly showed activation transients before lethargus but did not show many
731 transients during lethargus. RIM showed not only a change in transient frequency across
732 the lethargus cycle but also a reduction in baseline calcium activity. In *aptf-1(-)* mutant
733 worms, RIM continued showing calcium transients during lethargus, indicating that RIS
734 inhibits calcium transients in RIM during sleep bouts. However, reduction of baseline
735 calcium activity was preserved in *aptf-1(-)*, indicating that RIM activity is dampened
736 during lethargus independently of RIS at the level of baseline calcium activity. Together,
737 these experiments indicate that a dampening of behavioral and neural baseline activity that
738 is independent of RIS occurs during lethargus. This neuronal baseline and behavioral
739 dampening itself appears not to be sufficient to constitute normal sleep bouts but could
740 hypothetically lead to an activity change and decreased mutual inhibition in command
741 interneurons, thus promoting sleep induction [55,70].

742

743 **Fig 7. The dampening of neural and behavioral baseline activity levels during**
744 **lethargus is independent of RIS function.** Reduction of command interneuron activity
745 levels during lethargus occurs in wild-type worms and *aptf-1(gk794)* mutants. In the
746 wild-type condition, activity levels are reduced to -0.16 ± 0.02 . In the mutant condition,

747 activity levels are reduced -0.08 ± 0.02 . $**p < 0.01$, Wilcoxon signed rank test (S1 Data,
748 Sheet 7A and 7B). $\Delta F/F$, fluorescence change over baseline; GCaMP, genetically
749 encoded calcium indicator; RIS, Ring Interneuron S;

750

751 **An arousing stimulus inhibits RIS through RIM**

752

753 Arousal plays a major role in inhibiting sleep, but the circuits that mediate the effect of
754 arousing stimuli on RIS inhibition are not well understood. We therefore studied the circuit
755 by which stimulation of a nociceptor, the ASH (Amphid neuron class name) neurons, leads
756 to a reverse escape response and inhibition of RIS [71]. We optogenetically stimulated
757 ASH using ReaChR and green light and followed RIS and RIM activities. ASH activation
758 led to a strong activation of the RIM neuron and triggered a backwards response as
759 previously described [35,71]. Simultaneously, RIS was inhibited (Fig 8A). RIM can inhibit
760 PVC through reverse interneurons that it synchronizes [49,72]. Furthermore, strong RIM
761 activation can inhibit RIS more directly. To test whether ASH indirectly inhibits RIS
762 through RIM, we ablated RIM genetically by expression of *egl-1* from the *tdc-1* promoter
763 [35,38] and repeated the optogenetic stimulation of ASH. In RIM-ablated L4 animals,
764 activation of ASH caused the opposite effect on RIS activity. Instead of inhibiting RIS,
765 ASH activated RIS, while it still increased behavioral activity (Fig 8B). Consistent with
766 our calcium imaging data, ASH stimulation after RIM ablation predominantly caused a
767 forward locomotion response (Fig 8C). There are 2 ways ASH might inhibit RIS through
768 RIM. One possibility is that arousal strongly activates reverse interneurons, thus inhibiting
769 forward PVC neurons and RIS during stimulation. Consistent with this idea, gentle tail
770 touch increased RIS activity more strongly when RIM was ablated (S13 Fig). Another
771 option is that RIM inhibits RIS directly through tyramine and FLP-18. Both circuits might
772 play together (Fig 8D). These results delineate a circuit model for how sensory stimulation
773 can control RIS activation.

774

775 **Fig 8. Arousing stimulation inhibits RIS and sleep through RIM. (A) ASH**
776 **depolarization in wild-type worms leads to RIS inhibition and RIM activation, sleep**
777 **suppression, and mobilization. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon signed**

778 rank test for GCaMP and speed, Fisher's exact test for sleep fraction (S1 Data, Sheet 8A,
779 B). (B) ASH depolarization in RIM-ablated worms leads to weaker sleep suppression,
780 mobilization, and RIS activation. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon signed
781 rank test for GCaMP and speed, Fisher's exact test for sleep fraction (S1 Data, Sheet 8A,
782 B). (C) The response direction following ASH activation in wild-type worms is
783 predominantly reverse, while in RIM-ablated worms it is predominantly forward. $***p <$
784 0.001 , Fisher's exact test (S1 Data, Sheet 8C). (D) A circuit model for RIS regulation
785 through arousal by ASH. Activating synaptic input is shown as green arrows, inhibitory
786 synaptic input is shown as red arrows, and gap junctions are indicated as black
787 connections. RIM could serve as a synchronizer of AVE and AVA to regulate PVC and
788 therefore RIS inhibition. Additionally, RIM could inhibit RIS directly. ASH, AVA, AVE,
789 PVC, RIM, RIS, neuron class names; $\Delta F/F$, fluorescence change over baseline; GCaMP,
790 genetically encoded calcium indicator; n.s., not significant;

791

792 **Discussion**

793

794 **A wake-active circuit that controls locomotion also controls sleep**

795

796 Optogenetic activation and inhibition showed how the activity of presynaptic neurons
797 affects RIS depolarization during developmental sleep. Several presynaptic neurons can
798 activate RIS. RIM appears to be a potent direct inhibitor when activated strongly but can
799 also act as an activator of RIS. Loss-of-function experiments showed that the command
800 circuit controls activation of RIS, with PVC presenting a key activator of RIS. PVC has
801 long been known to mediate the forward escape response by transmitting information from
802 posterior sensory neurons to activate AVB premotor neurons to trigger forward locomotion
803 [46,49,50]. Consistent with promoting the forward escape response, optogenetic activation
804 of PVC leads to an increase in forward movement [50,73] (Fig 2D). Reverse movement, in
805 turn, is mediated by AVA, AVE, and AVD command premotor interneurons, which
806 activate reverse motor neurons. Forward PVC and reverse AVA/AVE/AVD command
807 interneurons are presynaptic to and mutually inhibit each other, which ensures discrete
808 forward and reverse locomotion states analogous to a flip-flop switch [49,55,74].

809

810 Our finding that PVC and RIM neurons present key activators of RIS that act in concert
811 suggests a model for how RIS is controlled; it also provides a potential mechanism for
812 linking sleep induction to decreasing arousal and for homeostatically maintaining a series
813 of sleep bouts. According to this model, during conditions of high arousal, such as during
814 development outside of lethargus, larvae are constantly awake. The command interneuron
815 circuit cycles between forward and reverse states, leading to the activation of forward or
816 reverse motor programs, respectively [49,74,75]. PVC activation has been associated with
817 the activity of forward states, and RIM has mostly been associated with the activity of
818 reverse states. Because neither activation of only the PVC nor of the RIM neurons appears
819 to be sufficient for sleep induction, RIS should not be activated sufficiently to induce sleep
820 during either forward or reverse states. At the transition between forward and reverse states,
821 locomotion pauses can occur. It has been shown that, in adult worms, RIS shows activation
822 transients in the nerve ring during locomotion pauses. These calcium transients appear to
823 be much smaller compared with activation transients during sleep bouts that extend to the
824 cell soma. Locomotion pausing is reduced after RIS ablation, suggesting that weak RIS
825 activation promotes pausing [76].

826

827 Lethargus induces a modest dampening of neuronal baseline activity that is independent of
828 RIS and that includes the RIM neurons. The RIM neurons become less excitable, which
829 should reduce their inhibitory effects on RIS and instead favor their activating effects. PVC
830 becomes resistant to inhibition and more potent in its capacity to activate RIS. We
831 hypothesize that these shifts in the properties of the interneurons of the locomotion circuit
832 favor the activation of the RIS neuron. RIS activation appears to require concerted
833 activation from PVC and RIM neurons (a process that is perhaps aided by other locomotion
834 interneurons). Both PVC and RIM appear to depolarize prior to RIS activation, and both
835 types of neurons contribute to RIS depolarization. This suggests that RIS might be
836 activated when both PVC and RIM exert activating effects. Such an overlapping activating
837 effect of PVC and RIM on RIS would most likely occur at the transition from forward to
838 reverse locomotion states, where there could be an overlap of both forward and reverse
839 neuronal activities. This would suggest that both locomotion stop and sleep bouts might be

840 induced by locomotion control interneurons at the transition between forward and reverse
841 locomotion states. The difference between locomotion stop and a sleep bout would be that,
842 in the former, RIS would only be modestly activated, whereas in the latter, RIS would be
843 strongly activated (Fig 9). Consistent with this model, sleep bouts are typically induced at
844 the end of long forward movements, whereas the exit from the sleep, for instance caused
845 by a noxious stimulus, is often through a reverse movement [70,75,77]. Arousal
846 promotes reverse command interneuron activity and strong RIM activation that can inhibit
847 RIS. Locomotion control and periods of behavioral activity and rest are already present in
848 animals that do not have a nervous system. It has therefore been hypothesized that sleep
849 and sleep-active neurons evolved from systems controlling locomotion activity and rest
850 [8]. The finding that a sleep-active neuron can also act as a locomotion pause neuron [76]—
851 and the discovery presented here that the locomotion circuit controls the depolarization of
852 a sleep-active neuron—suggests that sleep-controlling circuits might have evolved from
853 locomotion-controlling circuits and therefore that locomotion quiescence and sleep could
854 be regarded as homologous behaviors.

855

856 **Fig 9. A circuit model for RIS activation through locomotion interneurons. (A)**
857 **Activating synaptic input is shown as green arrows, inhibitory synaptic input is shown as**
858 **red arrows, and gap junctions are indicated as black connections. Outside of lethargus,**
859 **the nervous system cycles between forward and reverse states. RIS is not activated**
860 **sufficiently to cause a sleep bout, neither during the forward state during which PVC is**
861 **active nor during the reversal state during which RIM is active. The locomotion circuit**
862 **activates RIS briefly to cause a locomotion pause at the transition from forward to reverse**
863 **movement. Speculatively, the circuit that controls RIS during sleep also controls RIS**
864 **during locomotion pauses. (B) During lethargus motion bouts, the nervous system still**
865 **cycles between forward and reverse states. Baseline activity and excitability in RIM are**
866 **reduced, and PVC becomes resistant to inhibition and more potent to activate RIS. These**
867 **changes in locomotor interneurons shift the balance to favor strong RIS activation and**
868 **induction of a sleep bout, a process that may involve simultaneous activation from**
869 **multiple neurons, including RIM and PVC. Such an overlap activation of RIS by**
870 **otherwise mutually exclusive neurons could occur at the transition from forward to**

871 reverse locomotion states. Perhaps, RIS activation and sleep could occur similarly at the
872 transition from reverse to forward locomotion states. AVA, AVE, PVC, RIM, RIS,
873 neuron class names;

874

875 Our model suggests that the sleep switch is tripartite and includes not only wake-active
876 wake-promoting neurons and inhibitory sleep-active sleep-promoting neurons but also
877 wake-active sleep-promoting neurons as mediators of switch flipping. This sleep switch
878 acts as an amplifier that can translate a modest reduction of arousal into a massive
879 shutdown of behavioral activity during sleep. Dampening of neural activity and altered
880 properties of wake-active sleep-promoting locomotion neurons independently of sleep-
881 active neurons could be interpreted as a neural equivalent of sleepiness that leads to an
882 increased propensity to activate sleep-active neurons and to induce sleep bouts.

883

884 Mutations that increase arousal and suppress sleep increase the activity of reversal neurons,
885 whereas conditions that decrease arousal decrease the activity of the reversal neurons and
886 therefore increase the amount of sleep [67,68,74]. Also, the ablation of reverse command
887 interneurons such as AVE reduces reversals and leads to ectopic quiescence, as well as
888 increases sleep [46,52] (and this study). According to our model, increasing arousal should
889 increase the activity of RIM and other reverse command interneurons and thus should
890 inhibit RIS. Conversely, reducing arousal could promote weaker RIM activation and PVC
891 activation that should shift the equilibrium to stronger RIS activation.

892

893 What causes the termination of sleep bouts? The RIS neuron might not be able to sustain
894 prolonged activity, leading to the spontaneous cessation of a sleep bout. The RIS activation
895 transient and thus sleep bout can be blunted prematurely by a sensory or optogenetic
896 arousing stimulus [25,27,70,78]. Arousing stimulation, for instance, by activating the
897 nociceptive sensory neurons, triggers a reverse escape response through backwards
898 command and RIM interneurons [35,72,75,79]. Strong optogenetic RIM depolarization
899 inhibits RIS, and stimulation of the nociceptive ASH neurons causes inhibition of RIS that
900 depends on RIM, suggesting that a main physiological role of strong RIM activation is to
901 inhibit sleep upon arousing stimulation, perhaps by synchronizing the reverse interneurons

902 [72]. RIM activation can inhibit sleep also in response to acute food deprivation [80,81].
903 Thus, RIM might present not only an activator of RIS but also an arousal module that can
904 be activated upon sensing various external conditions that signal the need to suppress sleep.

905

906 RIS inactivation leads to disinhibition of arousal and brain activity, starting anew the cycle
907 of locomotion interneuron activity and locomotion behavior. Depending on the arousal
908 levels, the locomotion circuit causes RIS reactivation and thus a return to sleep either
909 immediately or after a delay. The timing of the rebound activation can be controlled by the
910 level of arousal—with strong arousal leading to longer wake periods before the return to
911 sleep—whereas milder stimulations cause an immediate return to sleep [23]. Consistent
912 with this circuit model of recurrent RIS activation, RIS activity oscillates, resulting in the
913 typical pattern of sleep bouts that are interrupted by activity bouts [22]. This circuit design
914 allows homeostatic sleep maintenance of a series of consecutive sleep bouts with sensory
915 stimulation restarting the cycle of RIS activation, thus prompting an acutely increased RIS
916 activation causing the return to sleep (Fig 6A) [23,70]. Our model predicts that RIS calcium
917 transient strength is a function of prior behavioral activity. Consistent with this view, RIS
918 calcium transients are stronger at the beginning and end of lethargus, when motion bouts
919 are high, but are less pronounced in the middle of lethargus, when motion bouts are less
920 pronounced (Fig 1A) [22,23]. Thus, the tripartite flip-flop circuit design allows an
921 adaptation of RIS activity to the strength required to induce sleep bouts at a given
922 behavioral activity level.

923

924 Here, we have identified a circuit controlling sleep-active neuron depolarization in *C.*
925 *elegans*. This work built on the neural connectome and was facilitated by the small size
926 and invariance of the nervous system as well as the transparency of the organism. While
927 the *C. elegans* sleep circuit clearly is built from fewer cells than the mammalian sleep
928 circuit [8,82,83], there are many conceptual similarities. For instance, in both *C. elegans*
929 and humans, sleep is controlled by inhibitory sleep-active sleep-promoting neurons that
930 depolarize at sleep onset to actively induce sleep by inhibiting wake circuits. A main
931 difference is that humans have many brain centers each consisting of thousands of sleep-
932 active neurons [12]. The single RIS neuron is the major inhibitory sleep-active neuron

933 required for sleep induction in *C. elegans* [25]. Work in mammals revealed the general
934 principles of wake-active wake-promoting neurons and sleep-active sleep-promoting
935 neurons as well as their mutual inhibition. While this information explains the flip-flop
936 nature of sleep and wake states, there is no satisfactory understanding of what flips the
937 sleep switch, i.e., how wakefulness is detected when the system is set to sleep, prompting
938 the activation of inhibitory sleep-active neurons [6]. Our model for the operation of the *C.*
939 *elegans* sleep circuit indicates that flipping of the sleep switch can be understood if wake-
940 active sleep-promoting neurons are added to the switch model. In this tripartite flip-flop
941 sleep switch model, the sleep-active sleep-promoting center is activated by wake-active
942 neurons. This activation should, however, only occur when the system is set to sleep, a
943 state that could present a neural correlate of sleepiness.

944

945 Sleep is reversible by stimulation, and hyperarousal is the major cause for insomnia in
946 humans [3,84,85]. Homeostatic sleep maintenance is an essential feature of sleep and is
947 found from worms to humans [19-21,23]. R2 ring neurons in *Drosophila* present an
948 integrator of wake time, causing subsequently increased depolarization of dFB sleep-
949 inducing neurons, thus forming a chronic sleep homeostat [86,87]. In vertebrates,
950 serotonergic raphe neurons are active during wakefulness and can reduce behavioral
951 activity and increase sleep pressure [88]. Our model of a tripartite flip-flop circuit suggests
952 that wake-active sleep-promoting neurons are an essential part of an acute sleep homeostat
953 that translates acute brain activity into increased sleep neuron activity when the system is
954 set to sleep. Wake-active sleep-promoting neurons measure systemic activity, i.e., they
955 become active together with a global brain activity increase and can then activate inhibitory
956 sleep-active neurons. Thus, the interplay of sleep-active sleep-promoting and wake-active
957 sleep-promoting neurons form an oscillator that periodically sends out sleep-inducing
958 pulses. Macroscopically, sleep in mammals exists as cortical oscillations of global down
959 states, known as slow waves [89]. Micro-arousals trigger cortical up states that are followed
960 by cortical down states, known as k-complexes [19-21]. Both slow-wave activity as well
961 as k-complexes could be hypothetically generated by wake-active sleep-promoting
962 neurons.

963

964 **Materials and methods**

965

966 **Worm maintenance and strains**

967 *C. elegans* worms were grown on Nematode Growth Medium (NGM) plates seeded with
968 *Escherichia coli* OP50 and were kept at 15 °C to 25 °C [90]. Crossed strains were
969 genotyped through Duplex PCR genotyping of single worms [91]. The primer sequences
970 that were used for Duplex PCR can be found subsequently. To confirm the presence of
971 transgenes after crossings, fluorescent markers were used. All strains and primers that were
972 used in this study can be found in S1 Text and S1 Table.

973

974 **Strain generation**

975

976 DNA constructs were cloned with the 3-fragments Gateway System (Invitrogen, Carlsbad,
977 CA) into pCG150 to generate new strains [92]. The *ArchT*, the *ReaChR*, and the *egl-1* genes
978 were expression optimized for *C. elegans* [93]. The *tdc-1::egl-1* transgene specifically
979 expresses the apoptosis-inducing protein EGL-1 in RIM and RIC. Therefore, RIM and RIC
980 are genetically ablated in worms carrying this transgene. The ablation is probably
981 incomplete in L1 worms. The *nmr-1::egl-1* transgene leads to the expression of *egl-1* in all
982 command interneurons causing their genetic ablation. Similar to the *tdc-1::egl-1* transgene,
983 ablation might be incomplete in L1 worms. In both lines, *egl-1* was co-expressed with
984 *mKate2*, which was used to verify the genetic ablations. Transgenic strains were generated
985 by microparticle bombardment or by microinjection. For microparticle bombardment, *unc-*
986 *119(ed3)* was used. The rescue of the *unc* phenotype was therefore used as a selection
987 marker [94,95]. The transgenes were backcrossed twice against N2 wild-type worms to
988 remove the *unc-119(ed3)* background. Extrachromosomal arrays were generated by DNA
989 microinjection. DNA was injected in wild-type, mutant, or transgenic worms. For injection,
990 DNA was prepared as follows: construct 30–100 ng/μl, co-injection marker 5–50 ng/μl,
991 and pCG150 up to a concentration of 100 ng/μl if required. Positive transformants were
992 selected according to the presence of co-injection markers. A table of all plasmids and a
993 list of all constructs that were generated for this study can be found in S2 Table and S2
994 Text.

995

996 **Generation of gene modifications using CRISPR**

997

998 The following allele was designed by us in silico and was generated by SunyBiotech.

999 Correctness of the alleles was verified by using Sanger sequencing.

1000

1001 PHX816: *flp-11(syb816 [SL2::mKate2::linker(GSGSG)::tetanustoxin_LC]) X*.

1002

1003 The coding sequences of tetanus toxin light chain and mKate2 were codon optimized and

1004 intronized as described previously and were synthesized [93]. The final sequence can be

1005 found in S3 Text.

1006

1007 **Imaging**

1008

1009 **Cameras and software.** All imaging experiments were conducted using either an iXon

1010 EMCCD (512 × 512 pixels) (Andor Technology Ltd., Belfast), an iXon Ultra EMCCD

1011 (1,024 × 1,024 pixels) (Andor Technology Ltd., Belfast), a Photometrics Prime 95B back-

1012 illuminated sCMOS camera (1,200 × 1,200 pixels) (Nikon, Tokyo), or a Nikon DS Qi2

1013 (4,908 × 3,264 pixels) (Nikon, Tokyo). For the iXon cameras, the EM Gain was set

1014 between 100 and 200. The exposure times used were between 5 and 30 ms. Andor IQ 2

1015 and 3 (Andor Technology Ltd., Belfast) and NIS Elements 5 (Nikon, Tokyo) were used for

1016 image acquisition.

1017

1018 **Illumination systems.** A standard 100-W halogen lamp together with an infrared filter

1019 (Semrock Brightline HC 785/62) (Idex Health and Science, New York) were used for

1020 differential interference contrast (DIC) microscopy or bright-field imaging. For calcium

1021 imaging and blue light stimulation, an LED illumination (CoolLED) with a 490-nm LED

1022 and standard GFP filter set (EGFP, Chroma) were used. Optogenetic stimulations and RFP

1023 imaging were performed with an LED illumination (CoolLED) with a 585-nm LED and

1024 standard TexasRed filter set (Chroma Technology Corp, Bellows Falls).

1025

1026 **Agarose microchamber imaging**

1027

1028 Long-term imaging experiments were conducted in agarose microchambers as previously
1029 described [96,97]. To summarize, a PDMS mold was used to cast box-shaped indentations
1030 in a hydrogel, which consisted of 3% or 5% agarose dissolved in S-Basal [98]. Two
1031 different sizes were used. We imaged L1 larvae in $190 \times 190 \times 15 \mu\text{m}$ microchambers, and
1032 L4 larvae were imaged in $370 \times 370 \times 25 \mu\text{m}$ microchambers. Depending on the
1033 developmental state of the worm that was imaged, either pretzel-stage eggs or L3 larvae
1034 were picked into the chambers with OP50 bacteria. Before imaging, worms were kept at
1035 either 20 °C or 25 °C.

1036

1037 For time-lapse calcium imaging experiments, L1 worms were filmed every 5 s (Figs 1A,
1038 3H-J, 4B-D, 5A, S8C-F and S11A Figs), every 8 s (Fig 7, S9A, S9C and S10 Figs), or
1039 every 10 s (Figs 3F and 4A) with DIC or bright-field imaging and widefield fluorescence.
1040 The DIC and bright-field light source was left on continuously, filtered through an infrared
1041 filter, and was blocked by a shutter during fluorescence image acquisition. LED
1042 illumination was triggered by the EMCCD camera using the TTL exposure output of the
1043 camera. An objective with 20× magnification, an LED with 480 nm (light intensity was
1044 between 0.15 and 2 mW/mm²), and EM gain of 100–200 was used. With the 20× objective
1045 and a 0.7 lens, 4 worms could be imaged simultaneously in one field. One to four fields
1046 could be filmed in parallel in one experiment. These image sequences gave measurable
1047 neuronal calcium transients and clear DIC or bright-field images to identify pumping or
1048 nonpumping phases.

1049

1050 During the continuous experiments in Figs 3F, 4B-D and 5A, only DIC or bright-field
1051 images were taken.

1052

1053 **AVA inhibition experiment**

1054

1055 NGM plates were prepared with histamine (HA; Sigma Aldrich, St. Louis, 10 mM) as
1056 previously described [54]. Young adult worms expressing a HA chloride channel in AVA

1057 and control worms were picked onto NGM HA plates the night before the experiments.
1058 The next morning, eggs together with *E. coli* bacteria from the NGM HA plates were
1059 picked into microfluidic chambers and DIC imaged as previously described [96,97].

1060

1061 **Optogenetic experiments**

1062

1063 Optogenetic experiments were either conducted in agarose microchambers as described
1064 previously, or the worms were immobilized. For immobilization experiments, the agarose
1065 was solved in S-Basal. We used the following 3 methods of immobilization for optogenetic
1066 experiments:

1067 1. Immobilization on a 3% agarose pad with 25 mM Levamisole (Sigma Aldrich, St. Louis)
1068 (S6 Fig)

1069 2. Immobilization on a 10% agarose pad with 0.1 μm Polybead microspheres
1070 (Polysciences, Warrington) [99] (Figs 1B/ SDQL, 1C/ SDQL, 3B-D, S1A/SDQL, S5A/
1071 SDQL, S4 and S8A Figs)

1072 3. Immobilization on a 10% agarose pad with 0.1 μm Polybead microspheres [99] and 25
1073 mM Levamisole (Figs 1B/PVC, 1C/PVC, 2A, 2C, 3B-D, 4F-G, S1A/PVC, S5A/PVC, S7A,
1074 S7C-D, S8B and S8H Figs)

1075

1076 Worms were imaged within 30 min of immobilization. A 100 \times oil objective was used for
1077 illumination and imaging in most experiments. For images in S6A-B Fig, a 1.5 lens was
1078 added (S6A and S6B Figs). The imaging in Fig 2A was done using a 40 \times objective.

1079

1080 ReaChR for neuronal depolarization or ArchT for hyperpolarization was utilized. For
1081 optogenetic stimulation, a 585-nm LED and a standard TexasRed filter set were used.

1082

1083 For optogenetic experiments with L1 larvae, either L4-stage worms or young adult worms
1084 were prepicked onto NGM plates with all-trans-retinal (ATR; Sigma Aldrich, St. Louis)
1085 and grown at 20 $^{\circ}\text{C}$ or 25 $^{\circ}\text{C}$. During the 2 d after exposure to ATR, pretzel-stage eggs or
1086 L1 worms were taken from this plate for optogenetic experiments. For optogenetic
1087 experiments with L4 larvae, an agar chunk containing a mixed population of growing

1088 worms was added to NGM plates containing ATR. Worms for optogenetic experiments
1089 were taken from this plate within the next 2 d.

1090

1091 Calcium imaging was conducted with an interval of 3 s and with an exposure time of 5–
1092 200 ms. A standard optogenetic protocol included calcium imaging during a baseline. This
1093 was followed by a stimulation time, in which the worms were optogenetically stimulated.
1094 The 585-nm light exposure was continuous except for brief interruptions during the time
1095 calcium imaging was conducted. After the optogenetic stimulation, calcium images were
1096 acquired during a recovery period.

1097

1098 In mobile worms, this standard protocol was preceded by 20 DIC frames that were taken
1099 every 500 ms to determine whether the worm was pumping. The overall protocol was
1100 repeated every 15 to 30 min. L1 mobile worms were imaged with a 20× objective and a
1101 0.7 lens. Mobile L4 worms were imaged with either a 10× objective (Figs 8A-C) or a 20×
1102 objective (Fig 1B/CEP, 1C/URY, S1A/CEP, S5A/URY and S3 Figs). Fixed worms were
1103 usually imaged between 1 and 4 trials. A delay preceded the standard protocol to allow the
1104 worm to recover from immobilization and between trials. To specifically manipulate PVC
1105 and SDQL in Figs 1B/PVC, 1B/SDQL, 1C/PVC, 1C/SDQL, 2D, S1A/PVC, S1A/SDQL,
1106 S4, S5A/PVC, S5A/SDQL, and S7C-D Figs, the stimulating illumination was restricted to
1107 the neuronal areas. This was achieved by reducing the size of the field aperture of the
1108 fluorescence illumination. To activate a specific neuron, it was moved into the illuminated
1109 area by using an automated stage. To image RIS, this neuron was moved into the
1110 illuminated area by the automated stage, while the optogenetic light stimulus was switched
1111 off and imaging light was switched on. The details for optogenetic experiments can be
1112 found in S3 Table.

1113

1114 **Behavioral imaging during PVC activation**

1115

1116 Worms were prepared on retinal plates and picked into microchambers as described
1117 previously. A 20× objective was utilized for imaging. The entire chamber was imaged
1118 through bright-field imaging. For tail-specific illumination, the LED blend was adjusted to

1119 illuminate a circular area with a radius of 58 μm . The 580-nm LED was manually turned
1120 on after 1 min of imaging and off after 4 min of imaging. A Prior XY stage (Prior Scientific,
1121 Cambridge) was manually operated to keep specifically the tail of the worm in the by the
1122 LED-illuminated area during stimulation. Worms were imaged with a frame rate of 8 Hz.
1123 Only every eighth image was used for analysis.

1124

1125 **Activity measurements of command interneurons**

1126

1127 GCaMP3.3 was expressed in command interneurons using the *glr-1* promoter [100]. L1
1128 larvae were placed in microfluidic chambers and were imaged using a time-lapse protocol.
1129 One DIC and one GFP image was taken every 8 s using a 20 \times objective and a 0.7 lens. The
1130 490-nm intensity for GFP imaging was set to 0.15 mW/mm^2 . Intensity values of all
1131 command interneurons located in the head of worms were extracted manually and analyzed
1132 as one entity.

1133

1134 **Pan-neuronal activity measurements**

1135

1136 GCaMP6s and RFP were pan-neuronally expressed under the *rab-3* promoter [101]. As in
1137 the activity measurements of command interneurons, L1 lethargus was imaged in
1138 microfluidic devices. For the optogenetic experiment (Fig 6C), every 30 min, 20 DICs were
1139 taken first in order to determine lethargus. This was followed by GFP images that were
1140 taken all 5.8 s for 9 min. The 490-nm intensity was set to 0.07 mW/mm^2 . In the blue light
1141 stimulation experiment, additional RFP images were taken. A custom-written MATLAB
1142 code detected the mean intensity of all head neurons in each GFP and RFP frame. The head
1143 neurons were thus analyzed as one entity.

1144

1145 **Blue-light stimulation experiments**

1146

1147 L1 worms were placed in microfluidic chambers for blue light stimulation experiments.
1148 The protocol was repeated every 15 min. First, 20 DIC pictures were taken every 500 ms
1149 to determine whether the worm was pumping or not. Next, baseline GCaMP was imaged

1150 for 3 min, the stimulation phase then lasted 18 s, and a recovery phase was imaged for 3
1151 min. The 490-nm intensity for calcium imaging was 0.07 mW/mm². The 490-nm intensity
1152 for stimulation was set to 1.01 mW/mm² with a 20× objective. The same LED was used
1153 for calcium imaging and stimulation. The intensity levels were controlled with Andor IQ2
1154 software.

1155

1156 The RFP signal of the pan-neuronal strain was imaged in addition to the GCaMP signal
1157 during the protocol every 3 s with 585-nm LED illumination, which was set to 0.17
1158 mW/mm².

1159

1160 **Mechanical stimulation using dish tapping**

1161

1162 The mechanical tapping set up was described previously [67,102]. L1 larvae were imaged
1163 in microfluidic chambers using a 20× objective and a 0.7 lens. Microfluidic chambers were
1164 put in a specialized dish. The dish was tapped by a piston driven by an electromagnet. The
1165 piston and the electromagnet were held in a homemade aluminum frame as described
1166 previously [102] (model used was Kuhnke, product number H2246). The voltage used for
1167 stimulus application was 5 V; the tapping stimulus was applied between image acquisition
1168 using TTL triggering to avoid blurring. Imaging was controlled with Andor IQ2 software.
1169 The imaging protocol was repeated every 15 min. First, 20 DIC pictures were taken with a
1170 frequency of 2 pictures per second to determine the status of worms. Throughout all
1171 following steps, GCaMP measurements were taken every 3 s. The 490-nm intensity for
1172 calcium imaging was 0.15 mW/mm². Baseline GCaMP was measured over 3 min.
1173 Following the tap, GCaMP was measured for 3 min. This experiment was initially planned
1174 to be combined with optogenetic stimulation, therefore a weak green light stimulus was
1175 applied, starting 15 s before and ending 45 s after the tapping stimulation. The part of the
1176 experiment during which a green light stimulus was applied was selected for presentation
1177 in this study. Green light (585 nm) for stimulation was set to 0.17 mW/mm². Because we
1178 did not see any noticeable changes upon applying green light, we presume that it does not
1179 strongly affect the experiment.

1180

1181 **Simultaneous calcium imaging of RIS and PVC**

1182

1183 In order to simultaneously image RIS and PVC, L1 lethargus worms were transferred from
1184 a growing plate using a platinum wire worm pick and were fixed on 10% agarose pads with
1185 0.1 μm Polybead microspheres [99] and 25 mM Levamisole. The worms were then imaged
1186 through a 40 \times oil objective with an image taken every 3 s for 30 min with 490-nm light of
1187 1.35 mW/mm to image GCaMP. Fluorescence intensities for PVC and RIS were cropped
1188 by using a region of interest. A custom-written MATLAB script then detected all RIS
1189 peaks. For this, the GCaMP data were first smoothed over 30 values through the in-built
1190 function “smooth,” which is a first-degree polynomial local regression model. Through the
1191 in-built MATLAB function “islocalmax,” and a minimum prominence value of 0.2, the
1192 locations of RIS peaks were detected. The data of RIS as well as PVC GCaMP intensity
1193 were aligned to the detected RIS peak location.

1194

1195 **Spinning disc confocal microscopy**

1196

1197 L4 worms were fixed with Levamisole. Spinning disc imaging was done with an Andor
1198 Revolution disc system (Andor Technology Ltd., Belfast) using a 488-nm (0.34 mW/mm²)
1199 and a 565-nm (0.34 mW/mm²) laser and a Yokogawa (Japan) CSU-X1 spinning disc head.
1200 Worms were imaged through a 100 \times oil objective. In S6A-B Fig, an additional 1.5 lens
1201 was used. z-Stacks with z-planes 0.5 μm apart spanning a total distance of 10 μm were
1202 taken, and a maximum intensity projection was calculated in ImageJ (developed by Wayne
1203 Rasband, open source).

1204

1205 **Tail-touch experiment**

1206

1207 L4 worms were grown and filmed on NGM plates with OP50 bacteria at 20 °C. An eyelash
1208 was used to gently touch the tail of the worms during L4 lethargus. The time from tail touch
1209 until the worms were immobile again was measured with a timer. If worms did not mobilize
1210 upon tail touching, the time was counted as zero. For GCaMP intensities, worms were

1211 imaged before and after tail touch each second for a total of 30 s. They were illuminated
1212 with a Leica EL6000 LED (Leica, Wetzlar).

1213

1214 **Image analysis**

1215

1216 Image sequences for analysis were selected either based on lethargus or molting time
1217 points. Lethargus was determined through DIC or bright-field images as the nonpumping
1218 phase before molting. Time points were classified to be in or outside of lethargus.
1219 Typically, the entire lethargus time and 2 h before lethargus were analyzed. Worms that
1220 were immobilized during the measurements were classified according to their pumping
1221 behavior on NGM plates directly before imaging. Two parameters were extracted from the
1222 image sequences, as follows.

1223

1224 1. Calcium signals were extracted automatically or manually with custom-written
1225 MATLAB codes. These codes extracted defined regions of each image and detected
1226 intensity and position data. Extracted regions were chosen slightly bigger than the sizes of
1227 measured neurons. From these extracted regions, a certain percentage of highest-intensity
1228 pixel was taken as signal. The remaining pixels were taken as background. From the signal,
1229 the background was subtracted. For the pan-neuronal and interneuron activity
1230 measurements, the signal in the head was treated as one large neuron and analyzed in the
1231 same way as single neurons. All head neurons expressed under the *rab-3* promoter were
1232 included in the pan-neuronal GCaMP measurements.

1233

1234 For all stimulation experiments, optogenetic and blue light stimulation experiments, the
1235 baseline measurement of each time point was utilized for signal normalization and $\Delta F/F$
1236 generation, except for Fig 6C. In Fig 6C, a mean of all baseline intensities for all wake time
1237 points for each worm was calculated. The mean was then utilized for normalization for all
1238 time points for each worm to better show the different RIS activities during wake and sleep.
1239 The pan-neuronal signal in Fig 6B was normalized over the measured RFP signal to
1240 retrieve $\Delta R/R$. For the transient alignments in Fig 3G, peaks and corresponding speeds
1241 were extracted through a custom-written MATLAB script and aligned as time point zero.

1242

1243 2. The speeds of the worms were calculated from the positions of the tracked neuron, except
1244 for experiments in which no GCaMP intensity was measured. To analyze these
1245 experiments, frame subtraction of DIC or bright-field images was done with a custom-
1246 written MATLAB routine instead.

1247

1248 **Baseline extraction**

1249

1250 In S12A and C Fig, the baseline of RIM GCaMP data was extracted by excluding the 95th-
1251 to 100th-percentile range for wild type and by excluding the 75th- to 100th-percentile range
1252 for *aptf-1(gk794)* mutants. The baseline was smoothed through a second-degree
1253 polynomial local regression model and with weighted linear least squares. Zero weight was
1254 assigned to data points 6 means outside the absolute deviation. The number of data points
1255 used for smoothing was 3%.

1256

1257 **Sleep-bout analysis**

1258

1259 Sleep bouts were extracted from selected parts of the time-lapse movies. Dependent on the
1260 experiment, a specific period of the movie sequence was selected and processed:

- 1261 1) The lethargus period (Figs 3F-J, 4A, 5A, 7, S9A, S9C, S10E-I and S12 Figs)
- 1262 2) The period from 2 h before lethargus up to the end of lethargus (Fig 1A)
- 1263 3) Either 3 h (Fig 4B-D and S8C-F Fig) or 4 h (S10A-D Fig) before shedding of the
1264 cuticle

1265 To extract sleep bouts, speeds and subtraction values were first smoothed. In Figs 1A, 3G-
1266 J, 4A-D, 5A, S8C-F, S10, and S12 Figs, speeds were smoothed through a first-degree
1267 polynomial local regression model over 20 time points. Other experiments were smoothed
1268 through a second-degree polynomial local regression model and with weighted linear least
1269 squares. Zero weight was assigned to data points 6 means outside the absolute deviation.
1270 Data were smoothed either over 3% of all data (Figs 3H-J, 7 and S9C-F Fig) or over 40
1271 data points (S9A, S9C, S12A and S12C Figs). This was achieved with the “smooth”
1272 function in MATLAB. Smoothed speeds were normalized between 0 and 1, with 0

1273 representing the lowest and 1 the highest smoothed speed value of each worm. In order to
1274 be scored as a sleep bout, the normalized speed had to be under a defined percentage
1275 threshold of the normalized speed for a minimum time. The exact speed and time thresholds
1276 were adjusted empirically to represent the worms' behavior [103]. In Fig 3G-I and S8C-F
1277 Fig, worms had to have a speed below 5% of their maximum smoothed speeds for at least
1278 2 min in order to be counted as sleeping. For all other experiments, the speed threshold was
1279 10%, and the time threshold was 2 min. The 2-min time threshold was implemented to
1280 exclude short pauses of the worm that may not represent sleep bouts. It was determined
1281 empirically. The sleep-bout analysis was carried out with a custom-written MATLAB
1282 script.

1283

1284 For stimulation experiments, the baseline and recovery time measurements were too short
1285 to include a minimum time threshold in the sleep-bout analysis. Therefore, immobility was
1286 used as a proxy for sleep. A mean of the wake speeds was calculated for each worm.
1287 Depending on the strain used, the worms were counted as sleeping when they were below
1288 a threshold of 5% to 30% of their mean wake speed. In most experiments, a worm was
1289 counted as sleeping when its speed was below 10% of the calculated mean of the wake
1290 speeds. To account for different locomotor behavior of the worms, in S11B Fig, the
1291 threshold was adjusted to 5%; in S8G Fig, to 20%; in S7B Fig, to 25%; in Fig 6C, S1C,
1292 S2A-C, S3, S5B, S9D-E, and S11I Figs, to 30%; and in S11C-E Fig, to 50%. RIS signals
1293 and speeds of wild type and mutants were aligned to sleep-bout onset for comparison in
1294 Figs 3H, 4A, S8F, S9A, S10D, and S10H Figs. RIM signals and speeds were aligned to
1295 sleep-bout onset in Fig 4A. For GCaMP normalization, 10 data points before sleep-bout
1296 onset were taken as baseline in order to calculate $\Delta F/F$. In S11A Fig, motion bouts were
1297 assigned whenever there was no detected sleep bout.

1298

1299 **RIS peak alignment**

1300

1301 For RIS wide peak detection (Fig 3G-H), first the normalized GCaMP data were smoothed
1302 over 60 time points with the in-built MATLAB function "smooth." Wide peaks were then
1303 detected with the in-built MATLAB function "findpeaks" and a minimum peak

1304 prominence threshold of 0.15. GCaMP intensities, speeds, and sleep fractions were then
1305 aligned to the detected peak maxima. Analysis for narrow peaks was conducted similarly;
1306 only 2 aspects were changed (S8C-F Fig). To find narrow peaks, smoothing was limited to
1307 only 5 time points, and a minimum peak prominence threshold was set to 0.2.

1308

1309 **Detection of direction of movement**

1310

1311 The direction of movement was analyzed with a custom-written MATLAB script. This
1312 MATLAB script took 2 points, the nose and the pharynx, to calculate the direction. For 2
1313 consecutive images, the distance of the nose in the first image to the pharynx in the first
1314 image was compared to the distance of the nose in the second image to the pharynx of the
1315 first image. If the distance increased, the worm was counted as moving forward; if it
1316 decreased, it was counted as moving in reverse. If the worm was below a threshold of 2
1317 $\mu\text{m/s}$, it was counted as sleeping in experiments Figs 2D and 4E. The position of nose and
1318 pharynx were detected manually (Figs 2D and 4E). For correction of the stage movement
1319 while manually tracking PVC (Fig 2D), the position of a corner of the stage was used.

1320

1321 **Fitting**

1322

1323 The data in Fig 6D were fitted to an asymptote, and the data in Fig 6E were fitted to a
1324 BoxLucasFit1 with Origin software. The data in S11A Fig were fitted to a logistic
1325 regression using Origin software (OriginLab Corporation, Northampton). Exact functions
1326 and R^2 values can be found in the respective Figures.

1327

1328 **Statistics**

1329 Sample sizes were determined empirically based on previous studies. If possible,
1330 experiments were carried out with internal controls. If this was not possible, control and
1331 experimental condition were alternated. Researchers were not blinded to the genotype for
1332 data analysis, as data analysis was performed by automated routines. Sample exclusion is
1333 described in the respective Methods sections. To compare GCaMP intensities and speeds
1334 of one sample group at different time points, the Wilcoxon signed rank test was utilized.

1335 The Fisher's exact test was used to compare the sleep fractions of one sample group at
1336 different time points. The entirety of the baseline was compared to the entirety of the
1337 stimulation period unless otherwise stated through significance bars. Data from different
1338 strains were compared with either the Kolmogorov-Smirnov test or the Welch test. The *p*-
1339 values can be taken from the respective Figure descriptions. Depicted in the graph is the
1340 mean \pm SEM unless otherwise stated. The box in the box plots represents the interquartile
1341 range with the median. The whiskers show the 10th- to 90th-percentile range, and the
1342 individual data points are plotted on top of the box.

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1344

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1350

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1352

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1727

1728

1729 **Supporting information**

1730

1731 **S1 Fig. Weak optogenetic RIM depolarization using the *gcy-13* promoter can induce**
1732 **RIS activation or inhibition.** (A) Control experiments. Optogenetic depolarization of
1733 RIS presynaptic neurons without the addition of ATR. For statistical calculations,
1734 baseline neural activities (0–0.95 min) were compared to neural activity levels during the
1735 stimulation period (1–1.95 min). * $p < 0.05$, ** $p < 0.01$, Wilcoxon signed rank test for
1736 GCaMP (S2 Data, Sheet S1A). (B) Optogenetic RIC depolarization induced an RIS
1737 activity increase outside of and during lethargus. An average of all responsive trials is
1738 shown in this figure. Trials were classified as responsive or nonresponsive. In responsive

1739 trials, an RIS activity increase correlated with the onset of the stimulation period. In
1740 nonresponsive trials, no change in RIS activity levels could be seen. “n” represents the
1741 number of animals tested, and “r” represents the number of trials. For statistical analysis,
1742 RIS baseline activity levels (0–0.95 min) were compared to activity levels during (1–1.95
1743 min) and after (2–2.95 min) the stimulation. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$,
1744 Wilcoxon signed rank test for GCaMP and speed, Fisher’s exact test for sleep fraction
1745 (S2 Data, Sheet S1B). (C) Depolarization of RIM using ReaChR expressed under the
1746 *gcy-13* promoter had no net effect on RIS function. Neural baseline activity levels (0–
1747 0.95 min) were compared to neuronal levels during the stimulation (1–1.95 min) and after
1748 the stimulation (2–2.95 min). * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, Wilcoxon signed rank
1749 test for GCaMP and speed, Fisher’s exact test for sleep fraction (S2 Data, Sheet S1C-E).
1750 (D) RIM optogenetic depolarization using ReaChR expressed under the *gcy-13* promoter
1751 induced either RIS activation or inhibition. Single trials were classified as activating if an
1752 activity increase in RIS correlated with onsets of optogenetic stimulation periods. Trials
1753 were classified as inhibitory if an activity decrease in RIS correlated with onsets of
1754 optogenetic stimulation periods. “n” represents the number of animals tested, and “r”
1755 represents the number of trials. For statistical testing, baseline neural activities (0–0.95
1756 min) were compared to neural activity levels during the stimulation period (1–1.55 min).
1757 * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, Wilcoxon signed rank test for GCaMP and speed,
1758 Fisher’s exact test for sleep fraction (S2 Data, Sheet S1C-E). (E) Percentage of RIS
1759 activation and inhibition following optogenetic RIM activation in different lethargus
1760 phases. Lethargus of each individual worm was split into 3 phases of comparable size
1761 (lethargus onset, middle of lethargus, and lethargus end). In each interval, for all worms
1762 tested the amount of trials showing an RIS activation or RIS inhibition were compared to
1763 the total amount of trials in this interval (S2 Data, Sheet S1C-E).

1764
1765 **S2 Fig. RIM inhibition of RIS requires tyramine and FLP-18.** Optogenetic RIM
1766 manipulations in these experiments were all performed with ReaChR expressed from the
1767 *tdc-1* promoter. (A) Optogenetic RIM depolarization in *flp-18(db99)* single mutants.
1768 Outside of lethargus, RIS inactivation caused by RIM optogenetic depolarization was
1769 reduced to 37% of wild-type inhibition levels. During lethargus in *flp-18(db99)* mutants,

1770 animal inhibition levels were only 25% of wild-type level. Neuronal activity levels before
1771 (0–0.95 min), during (1–1.95 min), and after (2.5–2.95 min) optogenetic RIM
1772 depolarization were compared. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon signed
1773 rank test for GCaMP and speed, Fisher’s exact test for sleep fraction (S2 Data, Sheet
1774 S2A). (B) Optogenetic RIM depolarization in *tdc-1(n3420)* single mutants. Outside of
1775 lethargus, optogenetic RIM depolarization in *tdc-1(n3420)* single mutants no longer
1776 induced changes in RIS activity levels. During lethargus, inhibition levels during the
1777 stimulation period only reached 40% of wild-type levels. Neuronal activity levels before
1778 (0–0.95 min), during (1–1.95 min), and after (2.5–2.95 min) optogenetic RIM
1779 depolarization were compared. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon signed
1780 rank test for GCaMP and speed, Fisher’s exact test for sleep fraction (S2 Data, Sheet
1781 S2B). (C) Optogenetic RIM depolarization in *flp-18(db99)* and *tdc-1(n3420)* double
1782 mutants had no effect on RIS function. Neuronal activity levels before (0–0.95 min),
1783 during (1–1.95 min), and after (2.5–2.95 min) optogenetic RIM depolarization were
1784 compared. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon signed rank test for GCaMP
1785 and speed, Fisher’s exact test for sleep fraction (S2 Data, Sheet S2C). (D) Quantification
1786 of inhibition strength. RIS activity levels during optogenetic RIM depolarization in *flp-*
1787 *18(db99)*, *tdc-1(n3420)* and *flp-18(db99)*, and *tdc-1(n3420)* double mutants were
1788 compared to wild-type levels. Wild-type data are depicted in Fig 1B, RIM panel.
1789 Inhibition strength was calculated by subtracting RIS activity levels during (1–1.95 min)
1790 the stimulation from activity levels before the stimulation (0–0.95 min). Samples were
1791 tested for normal distribution using the Shapiro-Wilk test. Wild type and mutants were
1792 compared with a Welch test. $***p < 0.001$ (S2 Data, Sheet S2D-E). (E) Quantification of
1793 RIS activity levels following RIM optogenetic depolarization. Activity levels in *flp-*
1794 *18(db99)*, *tdc-1(n3420)* and *flp-18(db99)*, and *tdc-1(n3420)* double mutants were
1795 compared to wild-type levels. Wild-type data are depicted in Fig 1B in the RIM panel.
1796 For statistical calculations, RIS activity levels before the stimulation (0–0.95 min) were
1797 subtracted from activity levels after the stimulation (2.5–2.95 min). Samples were tested
1798 for a normal distribution using the Saphiro-Wilk test. To compare genotypes, a Welch
1799 test was performed for all conditions, except for the comparison of activity levels
1800 between wild type and *tdc-1(n3420)* single mutants during lethargus. The *tdc-1(n3420)*

1801 data were not normally distributed, and thus a Kolmogorov-Smirnov test was used. *** p
1802 < 0.001 (S2 Data, Sheet S2D-E).

1803
1804 **S3 Fig. RIM activation of RIS requires glutamatergic signaling.** (A) RIM optogenetic
1805 depolarization using ReaChR expressed under the *gcy-13* promoter induced robust RIS
1806 activation in L4 larvae. In the L4 larvae, RIS activation by RIM optogenetic
1807 depolarization was more robust compared with the same experiment in L1 larvae. No trial
1808 selection was required. For statistical analysis, RIS baseline activity levels (0–0.95 min)
1809 were compared to activity levels during (1–1.95 min) and after (2–2.95 min) the
1810 stimulation. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, Wilcoxon signed rank test for GCaMP
1811 and speed, Fisher’s exact test for sleep fraction (S2 Data, Sheet S3A). (B) The activating
1812 input of RIM optogenetic depolarization on RIS was almost completely abolished in *eat-*
1813 *4(ky5)* mutants. For statistical analysis, RIS baseline activity levels (0–0.95 min) were
1814 compared to activity levels during (1–1.95 min) and after (2–2.95 min) the stimulation.
1815 * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, Wilcoxon signed rank test for GCaMP and speed,
1816 Fisher’s exact test for sleep fraction (S2 Data, Sheet S3B).

1817
1818 **S4 Fig. Activation of RIS by PVC is strongly enhanced during lethargus.**
1819 Optogenetic PVC depolarization in L2 larvae led to RIS activation outside of and during
1820 lethargus. RIS activation during lethargus was strongly enhanced. Plotted data represent
1821 the average over all experimental trials. Neural activity levels before the stimulation (0–
1822 0.95 min) were compared to activity levels during the stimulation (1–1.95 min). * $p <$
1823 0.05 , ** $p < 0.01$, Wilcoxon signed rank test (S2 Data, Sheet S4).

1824
1825 **S5 Fig. Optogenetic hyperpolarization experiments.** (A) Control experiments.
1826 Optogenetic hyperpolarization of RIS presynaptic neurons without the addition of ATR.
1827 For statistical calculations, baseline neural activities (0–0.95 min) were compared to
1828 neural activity levels during the stimulation period (1–1.95 min). * $p < 0.05$, ** $p < 0.01$,
1829 Wilcoxon signed rank test for GCaMP (S2 Data, Sheet S5A). (B) Hyperpolarization of
1830 RIM using ArchT expressed under the *gcy-13* promoter had no net effect on RIS
1831 function. Neural baseline activity levels (0–0.95 min) were compared to neuronal levels
1832 during the stimulation (1–1.95 min) and after the stimulation (2–2.95 min). * $p < 0.05$,

1833 $**p < 0.01$, $***p < 0.001$, Wilcoxon signed rank test for GCaMP and speed, Fisher's
1834 exact test for sleep fraction (S2 Data, Sheet S5B, C). (C) RIM optogenetic
1835 hyperpolarization using ArchT expressed under the *gcy-13* promoter caused a decrease in
1836 RIS activity levels in selected trials. Single trials were classified as activating if an
1837 activity increase in RIS occurred at the onset of the optogenetic stimulation period. Trials
1838 were classified as inhibitory if an activity decrease in RIS occurred at the onset of the
1839 optogenetic stimulation period. "n" represents the number of animals tested, and "r"
1840 represents the number of trials. For statistical calculations, neural baseline activity levels
1841 (0–0.95 min) were compared to levels during the stimulation period (1–1.75 min). $*p <$
1842 0.05 , $**p < 0.01$, $***p < 0.001$, Wilcoxon signed rank test for GCaMP and speed,
1843 Fisher's exact test for sleep fraction (S2 Data, Sheet S5B, C). (D) Simultaneous
1844 optogenetic hyperpolarization of CEP and URY neurons does not induce changes in RIS
1845 activity levels. For statistical testing, baseline neural activities (2–2.95 min) were
1846 compared to neural activity levels during the stimulation period (3–3.95 min) and after
1847 the stimulation (6–6.95 min). $**p < 0.01$, $***p < 0.001$, Wilcoxon signed rank test for
1848 GCaMP and speed, Fisher's exact test for sleep fraction (S2 Data, Sheet S5D).

1849
1850 **S6 Fig. *zk673.11* is expressed in PVC, RID, and cholinergic motor neurons.** (A–B)
1851 Expression of *nmr-1* and *zk673.11* only overlaps in PVC in the tail. (C–D) Expression of
1852 *nmr-1* and *zk673.11* does not overlap in head neurons.

1853
1854 **S7 Fig. PVC has multiple functions.** (A) PVC excitability remained unchanged during
1855 lethargus. Experiments were performed in immobilized L1 larvae to ensure PVC-specific
1856 green light illumination. A long baseline of 10 min was used to achieve stable baseline
1857 conditions. Activity levels of PVC during optogenetic depolarization were
1858 indistinguishable outside and during lethargus. PVC displayed a negative rebound
1859 transient after optogenetic depolarization. However, there was no difference in the
1860 amount of negative rebound outside and during lethargus (S2 Data, Sheet S7A). (B) RIS
1861 showed a rebound after mechanical stimulation. This rebound was stronger in worms
1862 during lethargus, and only during lethargus was the RIS rebound accompanied by a
1863 strongly increased immobilization of worms. $**p < 0.01$, $***p < 0.001$, Wilcoxon signed
1864 rank test for GCaMP and speed, Fisher's exact test for sleep fraction (S2 Data, Sheet

1865 S7B). (C–D) Effects of PVC stimulation on AVB activity. L1 larvae were immobilized
1866 for optogenetic experiments to ensure cell-specific stimulation of PVC. AVB activated
1867 upon optogenetic PVC depolarization with the same response strength during and outside
1868 of lethargus. AVB displayed an oscillatory activity pattern in 44% of all trials in worms
1869 outside of lethargus. AVB activity oscillated in 70% of all trials during lethargus. $*p <$
1870 0.05 , $**p < 0.01$, Wilcoxon signed rank test for GCaMP (S2 Data, Sheet S7C-D).

1871

1872 **S8 Fig. Effects of optogenetic RIS activation and inhibition on PVC and RIM**

1873 **activity.** (A) RIS depolarizes during optogenetic activation in fixed animals. As controls,
1874 experiments were performed in the absence of ATR. $***p < 0.001$, Wilcoxon signed rank
1875 test (S2 Data, Sheet S8A). (B) RIS hyperpolarization led to a weak PVC depolarization
1876 outside and during lethargus. For statistical calculations, neural activities before the
1877 stimulation period (0–1 min) were compared to activity levels during the stimulation
1878 period (1–2 min). $*p < 0.05$, $**p < 0.01$, compared before and during stimulation,
1879 Wilcoxon signed rank test (S2 Data, Sheet S8B). (C–D) Sample trace of RIS activity and
1880 worm locomotion behavior 3 h before shedding of the cuticle of *aptf-1(gk794)* and *flp-*
1881 *11(tm2705)* mutants (S2 Data, Sheet S8C and S8D). (E–F) *flp-11(tm2705)* mutants have a
1882 significantly increased number of short RIS peaks that do not correlate with sleep. (E)
1883 $**p < 0.01$, $***p < 0.001$, Welch test. (F) $**p < 0.01$, Kolmogorov-Smirnov test (S1
1884 Data, Sheet 3G-I). (G) Optogenetic RIS depolarization has no effect on RIM activity
1885 outside of and during lethargus. Neuronal activity levels before (0–0.95 min) and during
1886 (1–1.95 min) the stimulation period were compared. $*p < 0.05$, $**p < 0.01$, $***p <$
1887 0.001 , Wilcoxon signed rank test for GCaMP and speed, Fisher’s exact test for sleep
1888 fraction (S2 Data, Sheet S8G). (H) Optogenetic RIS hyperpolarization induced increased
1889 RIM activity both outside of and during lethargus. Measurements were performed in
1890 immobilized L1 larvae to reduce measurement noise. Activity levels during baseline
1891 measurements (0–0.95 min) were compared to levels during optogenetic RIS
1892 manipulation (1–1.95 min). $*p < 0.05$, Wilcoxon signed rank test for GCaMP (S2 Data,
1893 Sheet S8H).

1894

1895

1896

1897 **S9 Fig. Command interneurons are required for RIS activation and sleep induction.**

1898 (A) RIS activation in sleep bouts was strongly reduced in command-interneuron-ablated
1899 worms. Samples were tested for normal distribution using the Saphiro-Wilk test. $*p <$
1900 0.05, Welch test (S2 Data, Sheet S9A-C). (B) Command-interneuron-ablated worms
1901 moved much slower than wild-type worms. Command interneurons were genetically
1902 ablated by expressing ICE from the *nmr-1* promoter. Samples were tested for normal
1903 distribution using the Saphiro-Wilk test. $***p < 0.001$, Welch test for the wake condition
1904 and Kolmogorov-Smirnov test for the sleep condition (S2 Data, Sheet S9A-C). (C)
1905 Sample traces of RIS activity levels and worm locomotion behaviors outside of and
1906 during lethargus in command-interneuron-ablated worms and wild-type worms. In
1907 command-interneuron-ablated worms, quiescence bouts occurred only around the middle
1908 of the lethargus period (S2 Data, Sheet S9A-C). (D–E) Mosaic analysis of worms
1909 expressing an extrachromosomal array of *nmr-1::ArchT*. Worms were selected that
1910 expressed the transgene only in head neurons (D) or head neurons and PVC (E).
1911 Neuronal activity levels before (2–2.95 min) and during (3–3.95 min) the stimulation
1912 period was compared. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon signed rank test
1913 for GCaMP and speed, Fisher's exact test for sleep fraction (S2 Data, Sheet S9D and
1914 S9E).

1915

1916 **S10 Fig. Glutamatergic signaling is required for sleep induction.** (A–D) Sleep-bout

1917 analysis of *eat-4(ky5)* mutant larvae. *eat-4(ky5)* animals lacked significant RIS activation
1918 at sleep-bout onset. Consistent with this finding, mutant worms displayed a strong
1919 reduction in quiescence during lethargus. Samples were tested for a normal distribution
1920 using the Saphiro-Wilk test. $**p < 0.01$, $***p < 0.001$, Welch test for comparisons of
1921 sleep-bout lengths, sleep-bout frequencies, and sleep fractions. Wilcoxon signed rank test
1922 for quantifications of RIS activity levels in sleep bouts (S2 Data, Sheet S10A-D). (E–I)
1923 Sleep-bout analysis of *nmr-1(ak4)* mutant animals. RIS activity levels in sleep bouts were
1924 slightly reduced in the mutant. *nmr-1(ak4)* mutants did not show a reduced amount of
1925 quiescence during lethargus. Samples were tested for a normal distribution using the
1926 Saphiro-Wilk test. $*p < 0.05$, Welch test for comparisons of sleep-bout frequencies, sleep

1927 fractions, maximum RIS activity levels in sleep bouts, and RIS activity levels at the end
1928 of sleep bouts. Kolmogorov-Smirnov test for the comparison of sleep-bout lengths (S2
1929 Data, Sheet S10E-I).

1930

1931 **S11 Fig. RIS rebound activation following optogenetic hyperpolarization requires**
1932 **synaptic transmission.** (A) RIS GCaMP transient intensities in wild-type worms are
1933 correlated with the length of the preceding motion bout. The longer the preceding motion
1934 bout, the stronger the RIS activation (S2 Data, Sheet S11A). (B) RIM was inhibited
1935 during and post hyperpolarization. $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon
1936 signed rank test for GCaMP and speed, Fisher's exact test for sleep fraction (S2 Data,
1937 Sheet S11B). (C) RIS was optogenetically hyperpolarized with stimuli lasting for 48 s
1938 (C), 5 min (D), or 10 min (E). Worms not showing a rebound activation transient were
1939 excluded from the analysis, which was no worm for 48 s-, 1 out of 7 worms for 5 min-,
1940 and 1 out of 13 worms for 10-min stimulation experiments. Data from these plots were
1941 used to generate a dose-response curve of optogenetic RIS hyperpolarization (Fig 6D-E).
1942 $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon signed rank test for GCaMP and speed,
1943 Fisher's exact test for sleep fraction (S1 Data, Sheet 6D,E). (F-H) Following optogenetic
1944 hyperpolarization, RIS displayed strong rebound activation during lethargus (F).
1945 Rebound activation was abolished in a strain that is deficient for neurotransmission
1946 specifically in RIS (*flp-11::TetX*). (G) Rebound activation was abolished also by a
1947 mutation that impaired global synaptic transmission (*unc-13(s69)*). (H) $*p < 0.05$, $**p <$
1948 0.01 , Wilcoxon signed rank test (S2 Data, Sheet S11F-H). (I) RIS showed a negative
1949 rebound following its own optogenetic depolarization. The strength of the negative
1950 rebound transient depended on the lethargus status of the worm. Worms during lethargus
1951 displayed a 3-times-stronger negative rebound compared to worms outside of lethargus.
1952 $*p < 0.05$, $**p < 0.01$, $***p < 0.001$, Wilcoxon signed rank test for GCaMP and speed,
1953 Fisher's exact test for sleep fraction (S2 Data, Sheet S11I).

1954

1955 **S12 Fig. RIM baseline activity levels are dampened during lethargus independently**
1956 **of RIS.** (A) Sample traces of RIM transient frequencies, RIM baseline activities, and
1957 worm locomotion behaviors outside of and during lethargus in wild-type worms and *aptf-*

1958 *l(gk794)* mutants (S2 Data, Sheet S12). (B) Wild-type worms, but not *aptf-1(gk794)*
1959 mutant worms, display changes in RIM transient frequencies across lethargus. Transient
1960 frequencies were assessed manually. To be counted as a transient, RIM activity levels
1961 had to be at least twice as high as baseline activity levels. *** $p < 0.001$ Kolmogorov-
1962 Smirnov test for wild-type condition, Welch test for mutant condition (S2 Data, Sheet
1963 S12). (C) The reduction of RIM baseline activity levels during lethargus is preserved in
1964 *aptf-1(gk794)* mutants. ** $p < 0.01$, Wilcoxon signed rank test (S2 Data, Sheet S12).

1965

1966 **S13 Fig. Assaying gentle tail touch reveals an inhibitory role of RIM on RIS.** (A)
1967 RIM ablation increases the reinstating of immobility following gentle tail touch during
1968 lethargus. * $p < 0.05$, Kolmogorov-Smirnov test (S2 Data, Sheet S13A). (B) RIM ablation
1969 increases RIS activation in response to gentle tail touch. ** $p < 0.01$. Kolmogorov-
1970 Smirnov test (S2 Data, Sheet S13B).

1971

1972 **S1 Text. A list of strains that were used during this study.**

1973

1974 **S2 Text. A list of generated constructs during this study.**

1975

1976 **S3 Text. Sequence of the strain PHX816, which was generated during this study.**

1977

1978 **S1 Table. List of primers that were used during this study.**

1979

1980 **S2 Table. List of plasmids that were used during this study.**

1981

1982 **S3 Table. Experimental details of all optogenetic experiments conducted during this**
1983 **study.**

1984

1985 **S1 Data. Raw data for all experiments from the main figures (Figs 1-9).**

1986

1987 **S2 Data. Raw data for the experiments from the supporting figures (S1-S13 Figs).**

1988