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Title

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Journal

Current Biology, 25(6)

Authors

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Publication Date

2015-03-16

DOI

10.1016/j.cub.2015.01.016

Peer reviewed



Published in final edited form as:

Curr Biol. 2015 March 16; 25(6): 702–712. doi:10.1016/j.cub.2015.01.016.

Sleep Interacts with A β to Modulate Intrinsic Neuronal Excitability

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SUMMARY

Background—Emerging data suggest an important relationship between sleep and Alzheimer’s Disease (AD), but how poor sleep promotes the development of AD remains unclear.

Results—Here, using a *Drosophila* model of AD, we provide evidence suggesting that changes in neuronal excitability underlie the effects of sleep loss on AD pathogenesis. β -amyloid (A β) accumulation leads to reduced and fragmented sleep, while chronic sleep deprivation increases A β burden. Moreover, enhancing sleep reduces A β deposition. Increasing neuronal excitability phenocopies the effects of reducing sleep on A β , and decreasing neuronal activity blocks the elevated A β accumulation induced by sleep deprivation. At the single neuron level, we find that chronic sleep deprivation, as well as A β expression, enhances intrinsic neuronal excitability. Importantly, these data reveal that sleep loss exacerbates A β -induced hyperexcitability and suggest that defects in specific K⁺ currents underlie the hyperexcitability caused by sleep loss and A β expression. Finally, we show that feeding levetiracetam, an anti-epileptic medication, to A β -expressing flies suppresses neuronal excitability and significantly prolongs their lifespan.

Conclusions—Our findings directly link sleep loss to changes in neuronal excitability and A β accumulation and further suggest that neuronal hyperexcitability is an important mediator of A β toxicity. Taken together, these data provide a mechanistic framework for a positive feedback loop, whereby sleep loss and neuronal excitation accelerate the accumulation of A β , a key pathogenic step in the development of AD.

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INTRODUCTION

Alzheimer's disease (AD) is the most common cause of dementia worldwide, whose burden, both in terms of human suffering and health care costs, is expected to rise sharply in the next few decades [1]. β -amyloid peptides ($A\beta$), which are generated from sequential cleavage of amyloid precursor protein (APP), have been strongly implicated as having a key role in the pathogenesis of AD by substantial histopathologic, biochemical, and genetic data [2]. Thus, there is intense interest in identifying modifiable factors that modulate $A\beta$. Emerging evidence suggests potentially important links between sleep and AD [3]. It has long been appreciated that patients with AD have impaired sleep/wake cycles, with fragmented and reduced sleep at night [4–6]. Similarly, mouse models of AD have been shown to exhibit reduced sleep during their consolidated period [7, 8]. In humans, β -amyloid deposition, as inferred by a decrease in $A\beta$ levels in cerebrospinal fluid, is associated with reduced sleep quality [9], and reduced sleep and poor quality sleep are associated with increased fibrillar $A\beta$ burden in the brain [10].

Intriguingly, recent data also support a bidirectional relationship between sleep and amyloid--i.e., not only may $A\beta$ accumulation impair sleep, but poor sleep may increase $A\beta$ burden [3]. In humans, consolidated sleep attenuates the risk of developing AD conferred by the *E4* allele of *Apolipoprotein E* (*ApoE4*, an important genetic polymorphism for AD) [11, 12]. Moreover, a prospective study revealed that markedly fragmented sleep, as measured by wrist actigraphy, increased the risk of developing AD by ~1.5-fold, compared to those with the least fragmented sleep [13]. Importantly, using mouse models of AD, Kang et al. (2009) demonstrated that chronic sleep deprivation led to an increase in $A\beta$ burden in the brain.

What are the mechanisms by which sleep could modulate $A\beta$, and thus impact AD? A recent study has suggested that the "glymphatic system"--a system of perivascular cerebrospinal fluid (CSF) channels and glial processes that serves to remove metabolic waste from neurons in the brain--is more active during sleep than in wake [14]. In this study, the authors also showed that radioactively labelled $A\beta$ injected into the cortex was cleared more efficiently during sleep, as compared to wakefulness. Another potential mechanism underlying the relationship between sleep and β -amyloid is an alteration in neuronal activity. Although the function(s) of sleep remain enigmatic, one proposed function of sleep is to downscale synaptic strength following the synaptic potentiation that occurs during wakefulness [15]. For example, in rodents, molecular and electrophysiological markers of synaptic strength are increased following wakefulness, as compared to following sleep [16]. $A\beta$ accumulation also appears to be dependent on neuronal activity. For example, $A\beta$ cleavage from APP is enhanced with increased neural activity [17–19].

The fruit fly *Drosophila melanogaster* has been shown to sleep [20–23], and also has been established as a model for AD [24, 25]. There are several fly AD models, and we focused on a model that uses direct expression of human $A\beta_{42}$ coupled to a signal peptide. $A\beta_{42}$ -expressing flies have been shown to recapitulate several key features of AD, including $A\beta$ deposition, age-dependent learning impairment, and neurodegeneration [26–28]. Here, using this model, we investigated the functional interactions between sleep, excitability, and $A\beta$.

Our findings support a bidirectional relationship between sleep and β -amyloid and argue that increased neuronal excitability is a key mechanism underlying the effects of sleep on $A\beta$.

RESULTS

$A\beta$ Expression Leads to Reduced and Fragmented Sleep

To investigate the effects of $A\beta$ expression on sleep behavior in *Drosophila*, we used an established model of AD, whereby $A\beta$ 40, $A\beta$ 42, or $A\beta$ Arctic peptides are expressed pan-neuronally [28]. $A\beta$ 40 is less aggregate-prone than $A\beta$ 42, whereas $A\beta$ Arctic carries a disease-causing mutation that leads to enhanced aggregation [28, 29]. In order to bypass developmental effects, we induced expression of these $A\beta$ peptides in all neurons during adulthood using *daughterless-Geneswitch (da-GS)* [30]. We examined daytime and nighttime sleep amount of *da-GS>UAS-A β 40*, *da-GS>UAS-A β 42*, and *da-GS>UAS-A β Arctic* flies, and found that sleep amount was not significantly affected with overexpression of $A\beta$ 40 (Figures 1B and 1C). In contrast, nighttime sleep, but not daytime sleep, was significantly reduced with overexpression of $A\beta$ 42, while both daytime and nighttime sleep were significantly decreased with overexpression of $A\beta$ Arctic (Figures 1A–1C). These data suggest a “dose-dependent” relationship between $A\beta$ aggregation and sleep amount. We next examined the sleep architecture of these flies and found no significant effect of expression of $A\beta$ 40 and $A\beta$ 42 on nighttime sleep bout number or duration. However, inducing $A\beta$ Arctic expression resulted in fragmentation of nighttime sleep, as evidenced by an increase in sleep bout number and reduction in sleep bout duration during the night (Figures 1D and 1E). Consistent with these findings, the sleep of AD patients has previously been reported to be fragmented at night [5, 6]. Together, these data suggest that, as is the case in humans and mice, $A\beta$ expression in flies leads to reduced and fragmented sleep.

Nighttime Sleep Deprivation Increases Amyloid Burden

Previous work in mice has shown that chronic loss of sleep leads to an increase in $A\beta$ burden, raising the intriguing possibility that poor sleep may promote the pathogenesis of AD [31]. We thus addressed whether this phenomenon was conserved in other animals such as fruit flies. To reduce the time needed to visualize $A\beta$ deposits, we mainly focused on *UAS-A β Arctic* [28]. In addition, because one of the core features of AD is memory loss [1] and the mushroom bodies (MB) in *Drosophila* play a critical role in learning and memory [32], we expressed $A\beta$ Arctic peptides in the Kenyon cells (KC) of the MB using *OK107-Gal4*. Using mechanical deprivation, we subjected *OK107-Gal4>UAS-A β Arctic* flies to nighttime sleep deprivation for 1 week. As the use of chronic mechanical deprivation would increase locomotor activity and potentially result in “physical stress,” we also subjected flies to a similar deprivation paradigm during the daytime, to control for this potential confounder. As shown in Figures 2A and 2B, daytime sleep deprivation effectively reduced daytime sleep, but left nighttime sleep relatively intact, whereas nighttime sleep deprivation markedly reduced nighttime sleep and led to an increase in daytime sleep, likely reflecting “rebound sleep.” As expected, daily activity counts of *OK107-Gal4>UAS-A β Arctic* subjected to nighttime sleep deprivation were significantly increased, compared to those of flies not subjected to mechanical deprivation. However, there was no significant increase in locomotor activity in these flies when compared to those undergoing daytime sleep

deprivation (Figure 2C). To assess A β burden, we immunostained the brains of these flies using 6E10, a monoclonal antibody that detects an N-terminal epitope on the A β 42 peptide. Strikingly, confocal imaging of whole-mount brains demonstrated a significant increase in A β signal in *OK107-Gal4>UAS-A β Arctic* flies undergoing chronic nighttime sleep deprivation, compared to no deprivation or daytime sleep deprivation (Figures 2D and 2E). To reproduce these findings with a different MB-expressing driver, we examined *MB247-LexA>LexAop-A β Arctic* flies [33] and obtained similar results (Figure S1A). We asked whether these findings were specific for the MB, and found that nighttime sleep deprivation significantly increased A β Arctic accumulation in two other brain regions, the pars intercerebralis (using *OK107-Gal4*) and antennal lobes (using *NP1227-Gal4*) (Figure S1A). To examine whether these effects were specific to using *UAS-A β Arctic*, we repeated these experiments using *UAS-A β 42*. As shown in Figures S1C-E, the effects of nighttime and daytime sleep deprivation on sleep amount, daily activity, and A β burden were all recapitulated using *OK107-Gal4>UAS-A β 42* flies. We next assessed whether mechanical sleep deprivation would increase accumulation of an unrelated protein. As shown in Figure S1B, there was no increase in GFP signal when *OK107-Gal4>UAS-GFP* flies were subjected to nighttime sleep deprivation. Finally, we asked whether other manipulations that enhance cellular stress would affect A β Arctic or GFP accumulation. Neither chronic exposure to 31°C or 1mM paraquat affected A β Arctic or GFP burden in the MB KC (Figure S1F). Together, these data suggest that loss of nighttime sleep specifically leads to an increase in A β burden in a fly model of AD.

Genetic Manipulation of Sleep Modulates Amyloid Burden

To further investigate how changes in sleep affect A β accumulation, we used genetic approaches to reduce or increase sleep, instead of mechanical deprivation. To do this, we used 2 binary expression systems: the Gal4/UAS system to manipulate neurons that regulate sleep and the LexA/LexAop system to express A β Arctic in the MB. We previously demonstrated that activation of a subset of dopaminergic (DA) neurons using *TH-D4-Gal4* significantly reduces sleep amount [34]. Thus, we used *TH-D4-Gal4* to drive expression of dTrpA1, a heat-inducible cation channel [35] to activate DA neurons in flies expressing A β Arctic in the MB (*MB247-LexA>LexAop-A β Arctic*). As shown in Figure 3C, conditionally activating this subset of DA neurons resulted in a significant decrease in nighttime sleep, compared to controls. Similar to flies undergoing mechanical nighttime sleep deprivation, these flies exhibited a significant increase in A β accumulation, compared to controls (Figures 3A and 3D). It was previously shown that the ExF12 fan-shaped body (FB) neurons promote sleep [36, 37], and we recently identified a restricted Gal4 driver (*R72G06-Gal4*) from the Rubin/Janelia Farm collection that contains these cells (data not shown). Thus, in order to address whether increasing sleep would cause the opposite phenotype, i.e. a decrease in A β burden, we generated *R72G06-Gal4>UAS-dTrpA1, MB247-LexA>LexAop-A β Arctic* flies. As expected, conditional activation of FB neurons in these flies resulted in an increase in daytime and nighttime sleep (Figure 3C). Importantly, genetically increasing sleep in these flies decreased A β burden (Figures 3B and 3D). These data thus provide further evidence that sleep can modulate A β burden, and suggest that enhancing sleep can reduce A β pathology.

Manipulation of Neuronal Excitability Alters A β Accumulation

Previous work suggests that sleep deprivation increases neuronal excitability and synaptic transmission [15, 38]. We thus investigated whether alterations in neuronal excitability might underlie the changes in A β burden that we observe with manipulations of sleep. We first asked whether increasing neuronal excitability would result in an increase in A β accumulation. Expression of the bacterial sodium channel NaChBac [39] in the MB along with A β Arctic (*OK107-Gal4>UAS-NaChBac, UAS-A β Arctic*) resulted in a significant increase in A β signal in the MB (Figure S2B). However, this manipulation simultaneously reduced nighttime sleep in these flies (Figure S2A), making it difficult to disentangle whether activating these cells alone leads to an increase in A β burden. Therefore, to further assess this issue, we examined A β levels in the ExF12 FB neurons in *R72G06-Gal4>UAS-NaChBac, A β Arctic* flies. In flies expressing NaChBac and A β Arctic simultaneously in these ExF12 FB neurons, daytime sleep was increased compared to controls (Figure S2C), and a significant increase in A β burden was observed in the ExF12 cells, compared to controls (Figure S2D). These data thus dissociate changes in sleep from changes in excitability and suggest that enhancing neuronal excitability acts downstream of changes in sleep to increase A β burden.

To further address whether changes in excitability act downstream of sleep in modulating A β , we asked whether reducing neuronal excitability could suppress the increase in A β signal seen with sleep deprivation. We used dORK C2, an outward rectifying potassium channel [40], to decrease excitability of MB KC neurons. As a control, we used the non-conducting version of this channel, dORK NC. Reducing neuronal excitability in MB KC cells led to a trend towards an increase in daytime sleep in animals not undergoing sleep deprivation, while a marked reduction in nighttime sleep was observed in all animals undergoing sleep deprivation (Figure 4A). As shown in Figures 4B and 4C, electrically inhibiting the MB KC cells essentially blocked the increase in A β burden caused by sleep deprivation. These data suggest that hyperexcitability is necessary for sleep deprivation-dependent increases in A β accumulation.

Sleep Deprivation Increases Intrinsic Neuronal Excitability

In order to investigate the functional relationship between sleep, neuronal excitability, and A β , we decided to examine different neuronal cell groups. The circadian network in *Drosophila* consists of ~150 neurons comprised of different cell groups with distinct patterns of neuronal activity [41, 42]. The lateral groups (the ventrolateral and dorsolateral) are particularly suitable for the identification and imaging of individual cells. In addition, the ventrolateral neurons can be readily accessed for whole-cell patch clamp recordings, and the large ventrolateral neurons (l-LNvs) in particular have been shown to play a role in sleep/wake regulation [43–48]. Thus, the l-LNvs can be used to examine the relationship between sleep, neuronal activity, and A β accumulation in a single cell type. We used the *cry-Gal4* driver to manipulate the l-LNvs and the dorsolateral (LNd) groups (Figure 5A). We first expressed A β Arctic in these cells and investigated whether chronic sleep deprivation increased A β burden. As shown in Figures 5B and 5C, nighttime sleep deprivation caused an increase in A β signal in l-LNv cells, compared to controls, but not in LNd cells (which may reflect higher baseline A β expression in those cells—data not shown).

Our previous data suggest that changes in excitability act downstream of alterations in sleep to modulate A β levels. Indeed, previous studies have suggested that sleep “downscales” synaptic strength and that consequently, sleep deprivation leads to increased synaptic transmission [15]. Thus, we sought evidence that sleep deprivation could directly affect neuronal excitability. To address this question, we performed whole cell patch-clamp recordings of l-LNvs from animals with or without sleep deprivation and examined both the spontaneous action potential (AP) firing rate and evoked firing responses. As expected, nighttime sleep deprivation resulted in a significant decrease in nighttime sleep (Figure S3C). In order to isolate the l-LNv neurons from most excitatory and inhibitory inputs, we performed these recordings in the presence of mecamylamine (50 μ M) and picrotoxin (250 μ M). As shown in Figures 5D and 5E, the spontaneous AP firing rate of the l-LNv neurons was significantly increased (~1.8-fold) in flies subjected to sleep deprivation vs controls. Similar data were also obtained from measuring evoked responses from these neurons. Evoked AP firing rate was increased at all measured depolarizing currents (Figure 5F), and the frequency–current (f – I) slope was significantly increased in sleep-deprived animals, compared with controls (Figure 5G). We did not observe any significant change in resting membrane potential (RMP) in sleep-deprived animals vs controls (Figures S3A and S3B). Together, these data suggest that sleep deprivation enhances both spontaneous and evoked measures of intrinsic neuronal excitability.

A β -Dependent Hyperexcitability is Exacerbated by Sleep Deprivation

Next, we examined whether A β expression itself could also modulate neuronal excitability. Work in mammals has been conflicting, suggesting that A β can cause either a decrease or an increase in neuronal excitability [49–51]. To address this question in the *Drosophila* brain, we performed whole-cell patch clamp recordings on the l-LNv neurons in flies overexpressing A β 40, A β 42, or A β Arctic using *cry-Gal4*. As shown in Figure 6A, expression of A β Arctic, but not A β 40 or A β 42, led to a significant increase in spontaneous AP firing rate compared to controls. We next examined evoked parameters of excitability and found that overexpression of A β Arctic, but not A β 40 or A β 42, resulted in a significant increase in intrinsic neuronal excitability. In particular, the frequency of AP firing was increased in response to a range of injected currents, and the f – I slope was similarly increased (Figures 6B and 6C). RMP was not significantly affected in any of these conditions (Figure S4A). Together, these data suggest that A β Arctic expression on its own can enhance intrinsic neuronal excitability.

Given our previous finding that sleep deprivation alone could cause neuronal hyperexcitability, we next asked whether chronic sleep deprivation would further exacerbate A β Arctic-induced hyperexcitability. We examined *cry-Gal4*>*UAS-A β Arctic* flies subjected to chronic sleep deprivation (Figure S4C) and found that these flies exhibited a further increase in spontaneous and evoked intrinsic neuronal excitability, compared to non-sleep deprived *cry-Gal4*>*UAS-A β Arctic* as well as controls (Figures 6D–6G). RMP was depolarized, further suggesting that excitability was increased in the presence of A β Arctic with sleep deprivation (Figure S4B). In summary, these data suggest that sleep deprivation exacerbates the intrinsic neuronal hyperexcitability induced by A β expression.

Sleep Deprivation and A β Expression Lead to Impairment of Specific K⁺ Channel Currents

Changes in excitability could reflect alterations in a variety of ionic currents. Given that K⁺ channels in particular have been implicated in the regulation of sleep previously, we focused on changes to K⁺ currents [52–57]. To identify possible mechanisms for the hyperexcitability observed in l-LNV cells following chronic sleep deprivation (Figure S4D), we recorded steady state activation of three types of K⁺ currents: A-type K⁺ current (I_A), sustained K⁺ current ($I_{K(V)}$), and Ca²⁺-activated K⁺ currents (K_{Ca}) under voltage-clamp configuration. We found that over a range of membrane potentials, all three currents from sleep-deprived flies were significantly reduced compared with non-deprived control flies (Figures S5A–S5C). However, at a potential (–30 mV) near the spike threshold, only K_{Ca} currents showed a significant reduction compared to non-sleep deprived controls (Figures 6J and S5F). We next subjected *cry-Gal4>UAS-A β Arctic* flies to sleep deprivation and found that, in addition to K_{Ca} currents, I_A currents became markedly reduced at potentials near the spike threshold (Figures 6H and S5D). In sleep-deprived animals with or without A β Arctic expression, there was no significant effect on $I_{K(V)}$ currents (Figures 6I and S5E). These data suggest that alterations in K_{Ca} and I_A currents may play a role in the changes of excitability observed in l-LNV neurons under conditions of sleep deprivation and A β Arctic expression.

Levetiracetam Prolongs the Lifespan of A β -expressing Animals

Our data point towards changes in neuronal excitability as a key mediator of the effects of sleep loss on A β deposition. Furthermore, A β expression itself can induce neuronal hyperexcitability. Interestingly, abnormal elevations in neuronal activity can be seen in a variety of “pre-Alzheimer” states, including mild cognitive impairment (MCI) and ApoE4 carrier status [58, 59]. Furthermore, recent studies in humans with MCI and an AD mouse model have suggested that reducing neuronal excitability using the anticonvulsant levetiracetam (LEV) improves performance in learning and memory tests [60, 61]. Therefore, we wished to test the functional relevance of the hyperexcitability observed in A β Arctic-expressing flies by using LEV. We first examined whether feeding LEV to A β Arctic-expressing flies would suppress the increased AP firing rate seen in their l-LNV neurons. As shown in Figures 7A and 7B, when *cry-Gal4>UAS-A β Arctic* flies were chronically fed LEV (5 mg/kg) in their food, the increased AP firing rate of l-LNV neurons was reduced back to control levels. Neuronal activity can be associated with changes in neuronal structure [62], and the LNV neurons have been shown to exhibit experience- and sleep-dependent morphological changes [63, 64]. Therefore, we asked whether A β Arctic expression coupled with sleep deprivation affected the synaptic morphology of these neurons and whether LEV might suppress this effect. The LNV neurons express the neuropeptide Pigment Dispersing Factor (PDF), and so we used anti-PDF to label the synaptic terminals of these cells. Sleep-deprived flies expressing A β Arctic in their LNV neurons displayed a significant increase in PDF⁺ puncta in the optic lobes, and LEV treatment significantly inhibited this effect (Figure S6A). However, given that PDF is a releasable neuropeptide, we cannot rule out that these changes reflect alterations in the production or release of PDF. We next asked whether LEV treatment could inhibit the increase in A β burden in l-LNV neurons seen with sleep deprivation. As shown in Figure

S6B, LEV treatment resulted in a trend towards suppression of the increased A β burden induced by sleep deprivation, although this effect was not statistically significant.

Finally, pan-neuronal expression of A β Arctic peptide in flies has previously been shown to reduce lifespan [28], and so we asked whether suppressing neuronal hyperactivity by feeding these flies LEV would prolong their lifespan. We examined lifespan in *elav-Gal4>UAS-A β Arctic* flies and found, as expected, that these flies exhibited a reduction in their median lifespan, when compared to *elav-Gal4* controls (60 vs 45 days for females and 61 vs 38 days for males) (Figure 7E). Strikingly, when *elav-Gal4>UAS-A β Arctic* flies were fed LEV (5 mg/kg), their median lifespan was significantly extended by ~16% and ~18% for females and males, respectively (Figures 7C–E). In contrast, LEV did not extend the lifespan of control *elav-Gal4/+* female or male flies, suggesting that these effects are specific to A β Arctic-expressing flies (Figures S6C and S6D). Together, these data strongly suggest that neuronal hyperexcitability is an important mediator of A β -induced toxicity.

DISCUSSION

Our study supports a bidirectional relationship between sleep and β -amyloid and points towards an intimate relationship between sleep, neuronal excitability, and A β . Loss of sleep leads to neuronal hyperexcitation, which in turn increases A β burden. A β expression both reduces sleep and further enhances neuronal excitability (Figure 7F). Furthermore, suppression of A β -induced neuronal hyperexcitability significantly prolongs lifespan, suggesting that abnormal neuronal activity is an important mediator of A β toxicity.

How are changes in sleep related to alterations in neuronal excitability? In animals ranging from flies to humans, sleep is associated with broad changes in patterns of electrical activity [65, 66]. Although the function of sleep remains controversial, one prominent hypothesis is that sleep functions to downscale synaptic strength [15]. Along these lines, prolonged wakefulness has been associated with an increase in evoked cortical local field potential amplitudes in rats [16] and transcranial magnetic stimulation (TMS) measures of cortical excitability in humans [67]. Our data support this hypothesis, as sleep deprivation in flies leads to hyperexcitability of I-LNv neurons. Another recent study found that the ExF12 neurons in *Drosophila* also exhibited increased excitability with sleep deprivation, although in that case, it was suggested that this phenotype was related to the specific sleep-promoting function of those neurons [37].

There are a number of studies on the effects of APP/A β expression on neuronal activity in animal models of AD [49]. These studies, which have largely been conducted in mammalian systems, have been conflicting, possibly because different cell types and neuronal networks may behave differently in response to exposure to A β . Here, we have recorded from a single cell type (the I-LNvs) from intact fly brains, and found that A β expression markedly increases intrinsic neuronal excitability, and that this effect is exacerbated by sleep deprivation. Interestingly, patients with epilepsy exhibit elevated amounts of β -amyloid plaque in their brains [68]. Moreover, seizures are commonly seen in patients with early-onset AD carrying mutations in Presenilin 1 [69], and TMS studies have found increased excitability of primary motor cortex in patients with AD [70].

Our study suggests that neuronal hyperexcitability is an important and early contributor to the pathogenesis of AD. Changes in neuronal excitability in AD likely predate neurodegenerative changes [71]. For example, in young mice overexpressing APP, hippocampal neurons were found to be hyperactive, prior to the formation of β -amyloid plaques [50]. Furthermore, recent evidence suggests that increases in neuronal excitability may be deleterious for cognitive function. Indeed, patients with amnesic mild cognitive impairment (MCI) exhibit increased hippocampal activation by high-resolution fMRI, and reduction of this hippocampal activation with levetiracetam improved memory performance in these subjects [60]. Similar observations have been made in a mouse model of AD and in aged rats with cognitive impairment [61, 72]. We now provide evidence that reducing the neuronal activity of A β -expressing flies with levetiracetam prolongs their lifespan. Thus, taken together, these findings imply that early treatment of preclinical AD patients with antiepileptic medications may be beneficial in slowing the course of disease. Our findings reveal an important interaction between A β and sleep loss in modulating neuronal excitability and suggest that sleep loss, neuronal hyperexcitability, and A β accumulation form a positive feedback loop. As therapeutic interventions exist to manipulate sleep as well as neuronal excitability, these data suggest that targeting these pathways may be a fruitful approach towards slowing the progression or delaying the onset of this incurable disease.

Experimental Procedures

Details of experimental procedures are available in the online Supplemental Information.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

We thank Drs. Tzumin Lee, Damian Crowther, Mark Stopfer, Paul Shaw, and the Bloomington Stock Center for fly stocks. We thank members of the Wu Lab, Tom Lloyd, and Marilyn Albert for helpful feedback. This work was funded by NINDS R01NS079584 (M.N.W.), a Burroughs-Wellcome Fund Career Award for Medical Scientists (M.N.W.), an Alzheimer's Association New Investigator Research Grant (M.N.W.), and a Synaptic Plasticity and Cognitive Disorders Award from the Brain Science Institute at Johns Hopkins (A.S. and M.N.W.).

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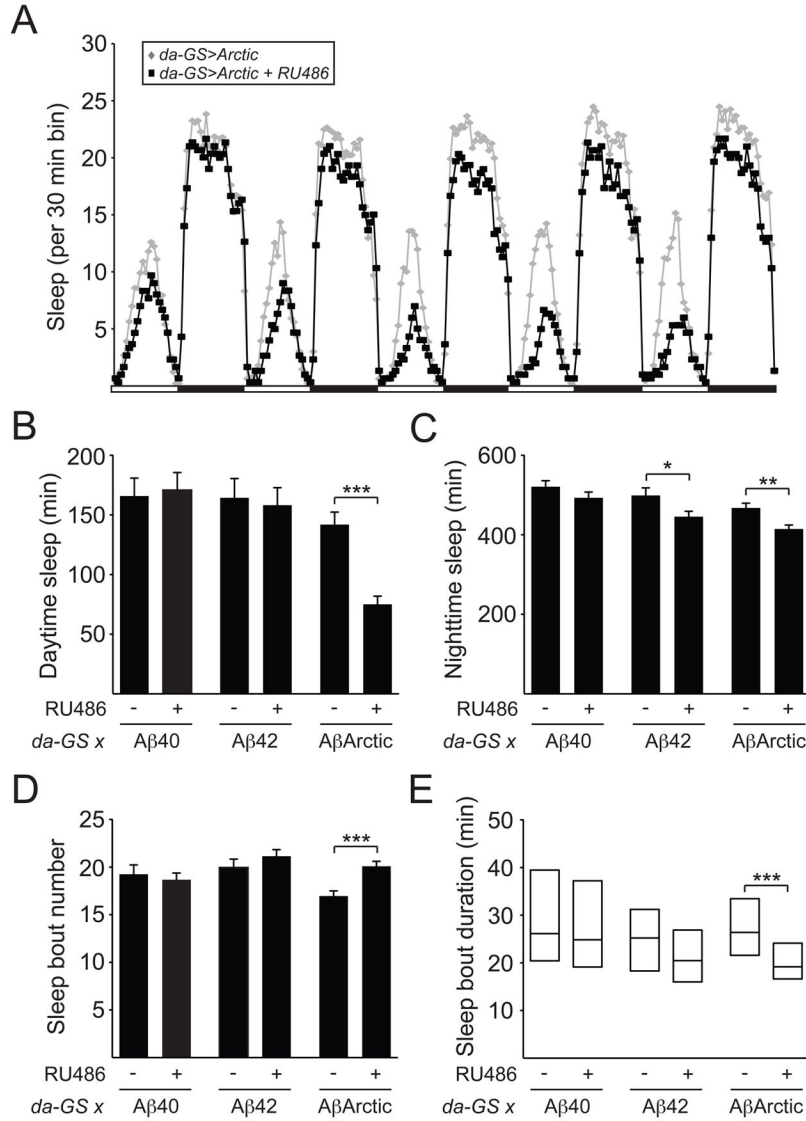


Figure 1. Induction of AβArctic expression reduces and fragments sleep
 (A) Sleep profile for *da-GS>UAS-AβArctic* flies fed 250 μM RU486 (black squares, n=92) or vehicle control (gray diamonds, n=77). Daytime sleep (B), nighttime sleep (C), nighttime sleep bout number (D), and nighttime sleep bout duration (E) for *da-GS>Aβ40* fed RU486 (n=48) or vehicle (n=38), *da-GS>Aβ42* fed RU486 (n=70) or vehicle (n=36), and *da-GS>AβArctic* fed RU486 or vehicle. Data in A are from the same flies as in B–E. In this and subsequent figures, error bars represent SEM. “*”, “***”, “****”, and “ns” denote $P < 0.05$, $P < 0.01$, $P < 0.001$, and not significant, respectively.

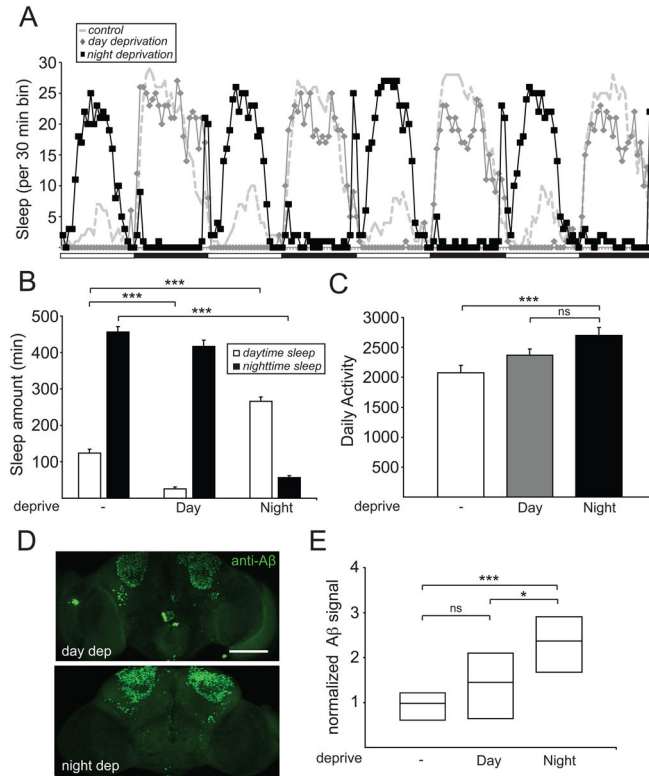


Figure 2. Mechanical sleep deprivation enhances A β burden

(A) Sleep profile for *OK107-Gal4>UAS-A β Arctic* flies undergoing no, daytime, or nighttime sleep deprivation from a representative experiment. White bars and black bars denote light and dark periods, respectively. Sleep amount (B) and daily activity (C) for *OK107-Gal4>UAS-A β Arctic* flies, where “-”, “Day”, and “Night” denote no, daytime, and nighttime sleep deprivation, respectively. (D) Representative whole-mount brain confocal images for *OK107-Gal4>UAS-A β Arctic* flies undergoing daytime (“day dep”) or nighttime (“night dep”) sleep deprivation, immunostained with anti-A β 42 antibody (6E10). Maximum projection images are shown. (E) Normalized A β signal intensity in the MB KC from *OK107-Gal4>UAS-A β Arctic* flies undergoing no (n=10), daytime (n=10), or nighttime (n=9) sleep deprivation. A β signal intensity is not normally distributed and is thus presented here and in subsequent figures as a simplified box plot with the median shown as the line inside the box, and the 75th and 25th percentiles shown as the top and bottom, respectively. Scale bar represents 100 μ m.

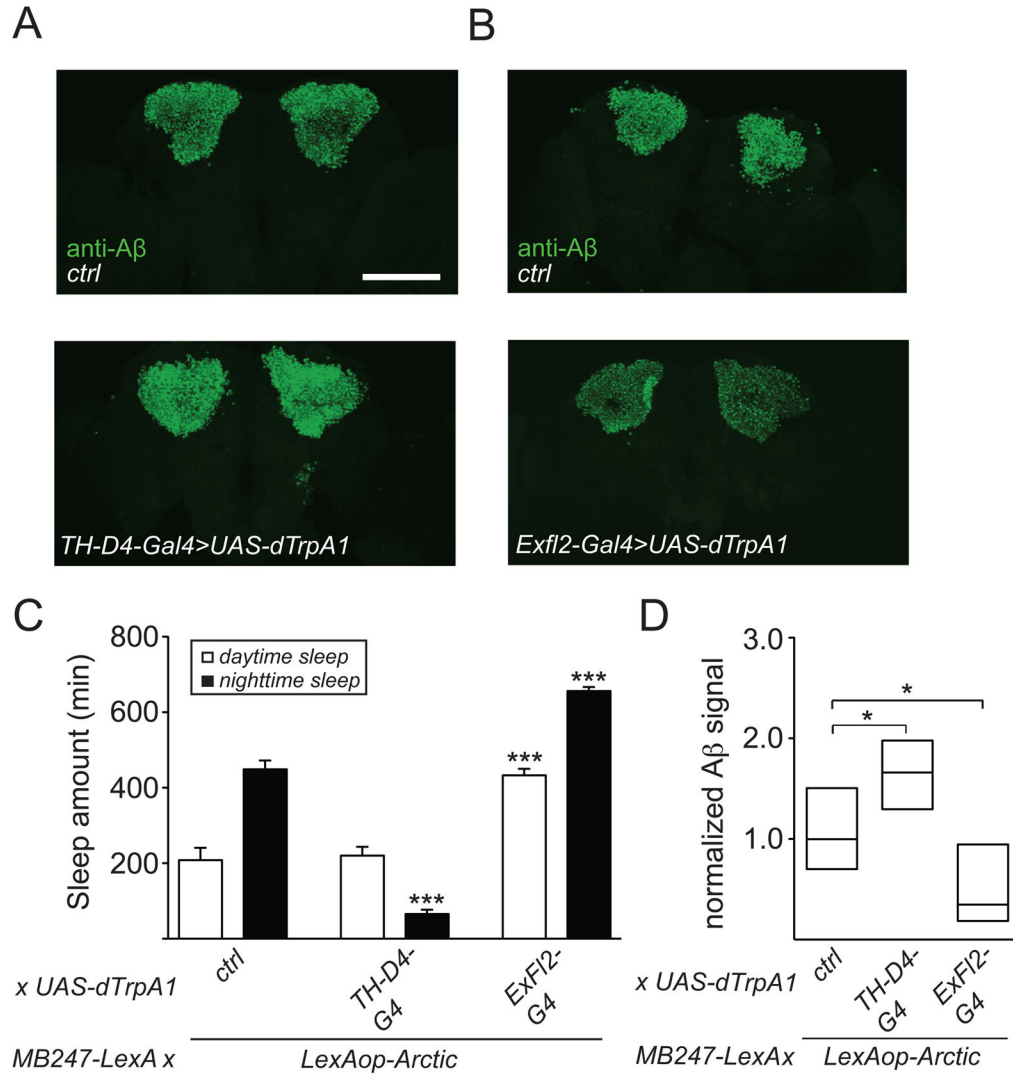


Figure 3. Genetic manipulation of sleep modulates Aβ levels
 (A and B) Representative maximum projection images of whole-mount brains immunostained with 6E10 from *LexAop-AβArctic/+; UAS-dTrpA1/MB247-LexA* (“ctrl,” top panels) and *LexAop-AβArctic/+; UAS-dTrpA1/MB247-LexA, TH-D4-Gal4* (“*TH-D4-Gal4*>*UAS-dTrpA1*,” bottom panel), and *LexAop-AβArctic/+; UAS-dTrpA1/R72G06-Gal4, MB247-LexA* (“*ExF12-Gal4*>*UAS-dTrpA1*,” bottom panel) flies. (C) Daytime and nighttime sleep amounts for *LexAop-AβArctic/+; UAS-dTrpA1/MB247-LexA* (n=24), *LexAop-AβArctic/+; UAS-dTrpA1/MB247-LexA, TH-D4-Gal4* (n=24), and *LexAop-AβArctic/+; UAS-dTrpA1/R72G06-Gal4, MB247-LexA* (n=22). (D) Normalized Aβ signal intensity in the MB KC for the flies in (C), shown as a simplified box plot. dTrpA1 was chronically activated by subjecting flies to 29°C for 1 week. Scale bar represents 100 μm.

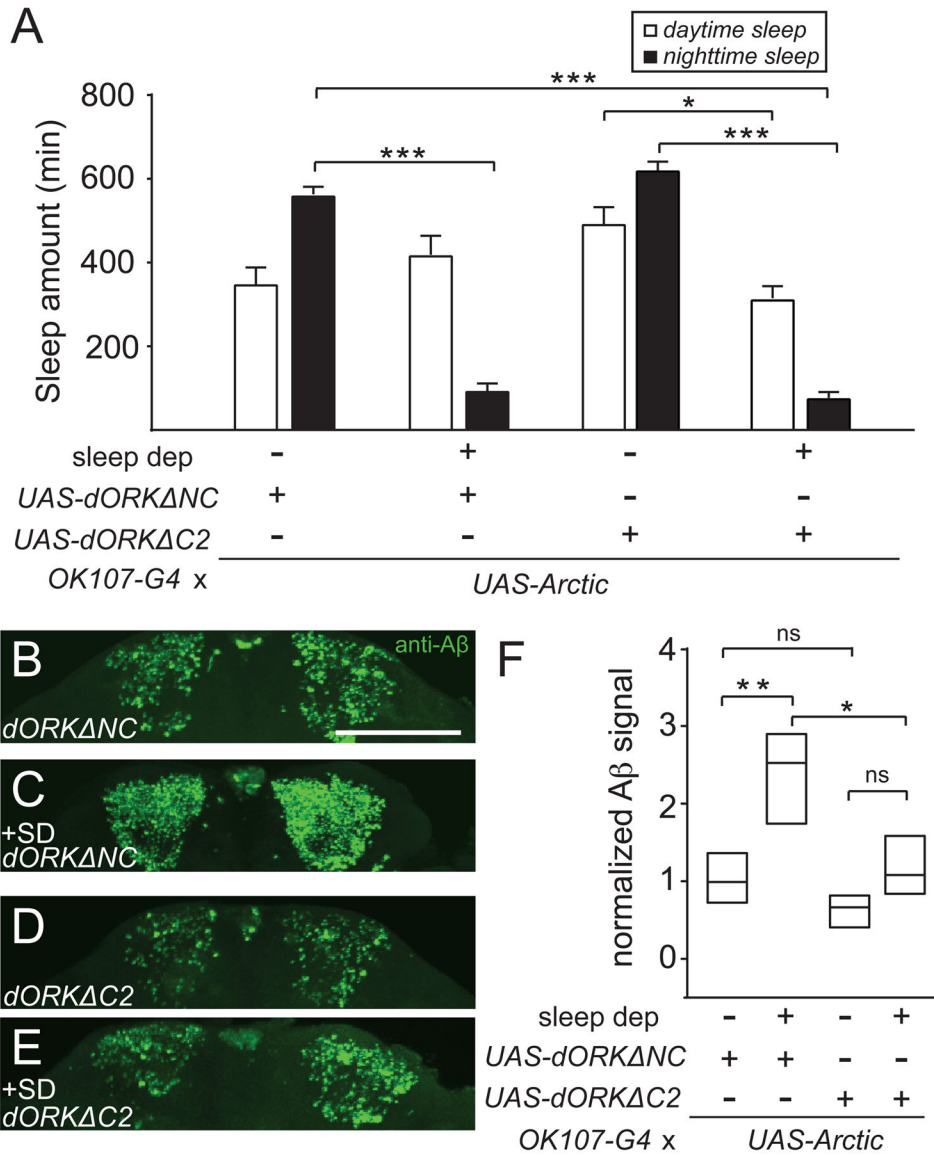


Figure 4. Inhibiting neuronal excitability suppresses A β accumulation induced by sleep loss
 Daytime and nighttime sleep amounts (A), representative maximum projection images of KC from whole-mount brains immunostained with 6E10 (B–E), and normalized A β signal intensity in the MB KC neurons (F) for *OK107-Gal4>UAS-A β Arctic*, *UAS-dORK NC* without sleep deprivation (n=20) and with sleep deprivation (n=13) and *OK107-Gal4>UAS-A β Arctic*, *UAS-dORK C2* without sleep deprivation (n=17) and with sleep deprivation (n=13). Scale bar represents 100 μ m.

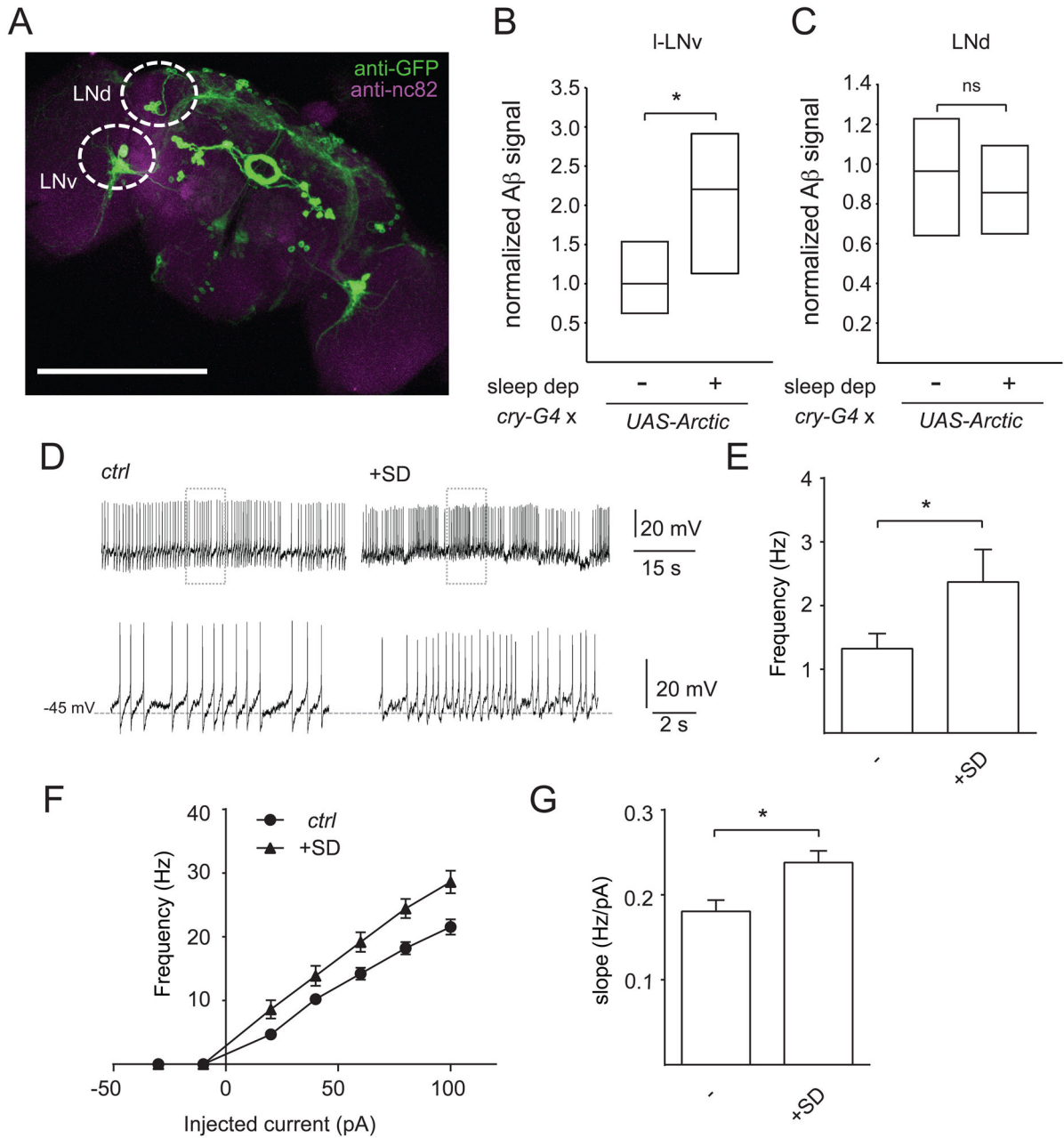


Figure 5. Sleep deprivation increases intrinsic neuronal excitability
 (A) Maximum projection of a whole-mount brain immunostained with anti-GFP from *cry-Gal4>UASCD8::GFP*. Normalized Aβ signal intensity in I-LNv (B) and LNd (C) cells for *cry-Gal4>UAS-AβArctic* with (n=24) or without sleep deprivation (n=21), shown as a simplified box plot. (D) Representative traces showing spontaneous AP firing of I-LNvs at ZT0-3 in *cry-Gal4>UAS-CD8::GFP* flies with or without sleep deprivation (SD). The bottom traces in (D) are expanded traces of the boxed regions in the top traces. Mean firing rate of spontaneous activity (E), mean frequency of spikes elicited in response to current injections with 300 ms stepping pulses at 20 pA increments, ranging from -30 pA to 100 pA

(F), and *f-I* slope (G) of l-LNV neurons in control (*cry-Gal4>UAS-CD8::GFP*) animals with (n=12) or without (n=15) sleep deprivation. Recordings were performed in the presence of mecamlamine (50 μ M) and picrotoxin (250 μ M), in order to isolate these cells from most excitatory and inhibitory inputs. Scale bar represents 200 μ m.

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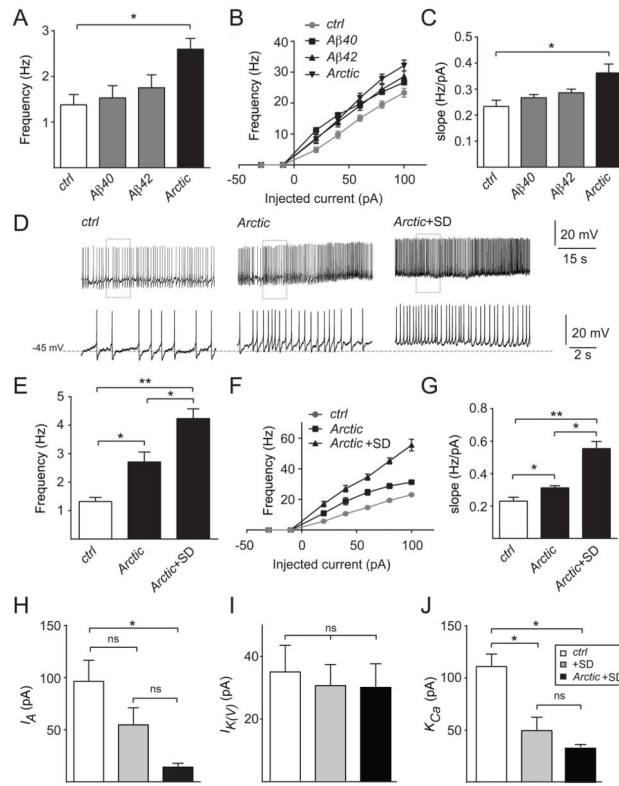


Figure 6. Sleep deprivation exacerbates $A\beta$ -dependent neuronal hyperexcitability

Mean firing rate of spontaneous activity (A), mean frequency of spikes elicited in response to current injections ranging from -30 pA to 100 pA (B), and *f-I* slope (C) of l-LNV neurons in control *cry-Gal4>UAS-CD8::GFP* ($n=16$), *cry-Gal4>UAS-A β 40, UAS-CD8::GFP* ($n=17$), *cry-Gal4>UAS-A β 42, UAS-CD8::GFP* ($n=15$), and *cry-Gal4>UAS-A β Arctic, UAS-CD8::GFP* ($n=18$). (D) Representative traces showing AP firing of l-LNV neurons in control vs *cry-Gal4>UAS-A β Arctic, UAS-CD8::GFP* flies +/- sleep deprivation (SD). Bottom traces in (D) are expanded traces of the boxed regions in the top traces. Mean firing rate of spontaneous activity (E), mean frequency of spike elicited in response to current injections ranging from -30 pA to 100 pA (F), and *f-I* slope (G) of l-LNV neurons in control ($n=17$) vs *cry-Gal4>UAS-A β Arctic, UAS-CD8::GFP* with ($n=15$) or without sleep deprivation ($n=19$). I_A (H), $I_{K(V)}$ (I), and K_{Ca} (J) current amplitude at the spike threshold (-30 mV) from l-LNVs for *cry-Gal4>UAS-CD8::GFP* with ($n=5, 10,$ and $11,$ respectively) or without ($n=5, 7,$ and $8,$ respectively) sleep deprivation and *cry-Gal4>UAS-A β Arctic, UAS-CD8::GFP* with sleep deprivation ($n=5, 6,$ and $4,$ respectively). Recordings were performed in the presence of mecaminylamine (50 μ M) and picrotoxin (250 μ M), in order to isolate these cells from most excitatory and inhibitory inputs.

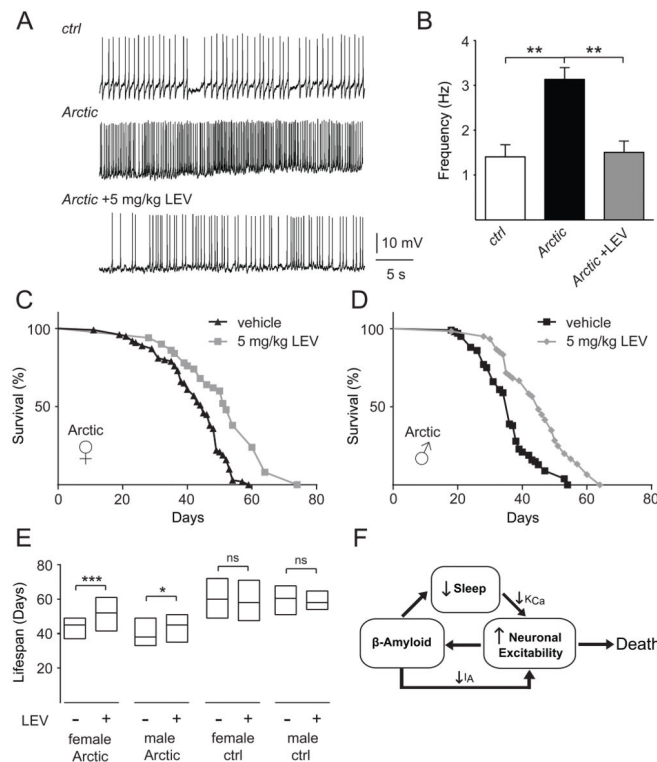


Figure 7. Levetiracetam suppresses neuronal firing and prolongs lifespan of A β Arctic-expressing flies

(A) Representative traces showing spontaneous firing of I-LNv neurons in control (*cry-Gal4>UASCD8::GFP*) vs *cry-Gal4>UAS-A β Arctic*, *UAS-CD8::GFP* flies fed vehicle or 5 mg/kg levetiracetam (LEV). (B) Quantification of mean firing rates shown in (A) ($n = 4$ for control, $n = 5$ for A β Arctic, and $n = 8$ for A β Arctic + LEV). Survivorship curves of *elav-Gal4>UAS-A β Arctic* female (C) and male (D) flies fed vehicle or 5 mg/kg LEV. (E) Lifespan extension of *elav-Gal4>UAS-A β Arctic* female and male flies by LEV, where lifespan is displayed as a simplified box-plot. Data for the *elav-Gal4>UAS-A β Arctic* (“Arctic”) flies shown here are the same as in (C) and (D) ($n = 98$ for vehicle- and $n = 52$ for LEV-fed females, and $n = 100$ for vehicle- and $n = 60$ for LEV-fed males). For *elav-Gal4/+* (“ctrl”) flies, $n = 30$ for vehicle- and LEV-fed females and males. (F) Model connecting sleep, neuronal excitability, and A β . Sleep loss leads to a reduction in Ca²⁺-dependent K⁺ currents, causing neuronal hyperexcitability. This enhanced excitability, in turn, results in increased A β accumulation. A β itself reduces sleep and further increases neuronal excitability via a decrease in voltage-gated K⁺ currents, generating a positive feedback loop whereby sleep loss and A β interact to substantially increase neuronal activity and A β burden. Increased neuronal excitability then contributes to reduced lifespan.