

# Elucidation of proteins controlling the sleep and awake transition

October 26 2022

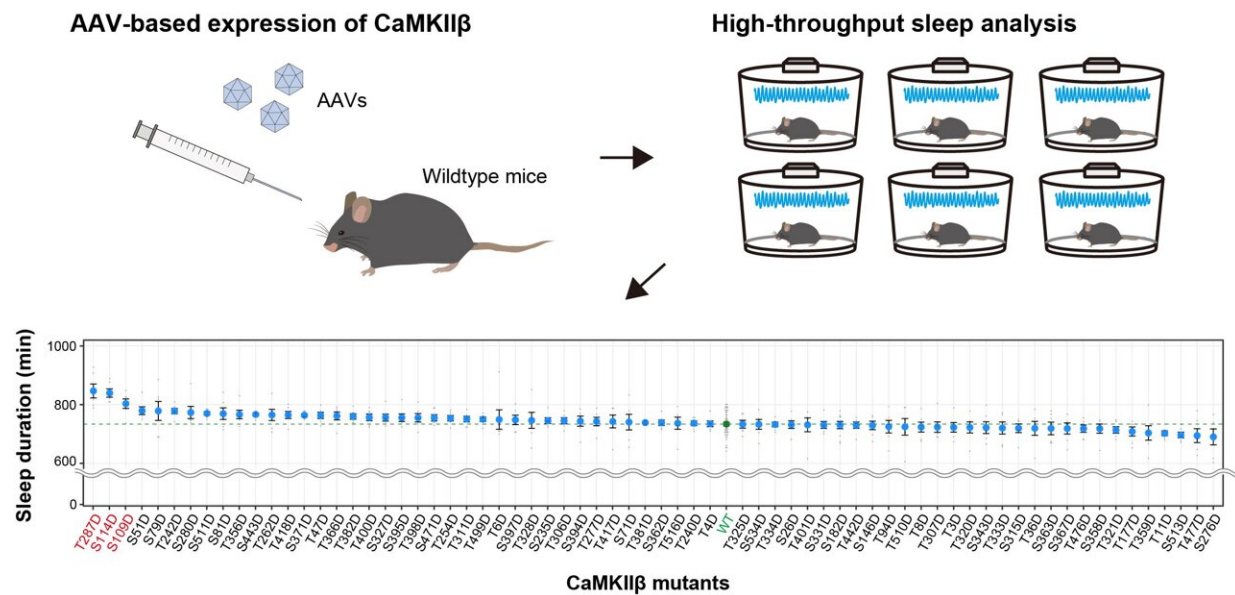


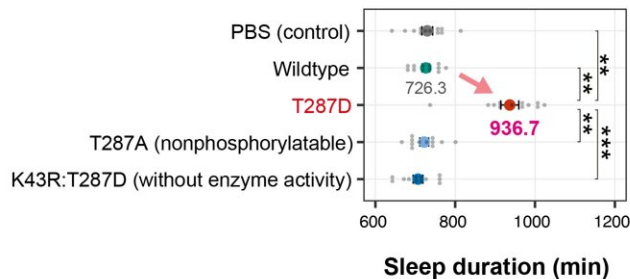
Fig.1 AAV-based screening for CaMKII $\beta$  phosphomimetic mutants Wild-type mice were administered with AAVs (adeno-associated viruses) to express the CaMKII mutant (upper right) and their sleep phenotypes were analyzed (upper left). Comprehensive analysis of mice administrated with 69 AAVs corresponding to CaMKII $\beta$  phosphomimetic mutants revealed that T287D (T287D), a phosphomimetic mutant of T287, remarkably extended sleep duration (below). Credit: Tone et al.

Professor Hiroki Ueda, Assistant Professor Daisuke Tone, Lecturer Koji

Ode, and Qianhui Zhang (4th-year graduate student at the time) at the University of Tokyo showed that CaMKII $\beta$ , a predominant protein kinase expressed in neurons, is involved in prolonging sleep duration via facilitating sleep onset and inhibiting awakening.

Several recent studies have shown that [protein phosphorylation](#) is important for sleep regulation. The aforementioned study group reported in 2016 that CaMKII $\alpha$  and CaMKII $\beta$  are [protein kinases](#) that promote sleep and suggested the possibility that protein phosphorylation is the likely molecular driver of somnolence and proposed phosphorylation hypothesis of sleep. CaMKII $\alpha$  and CaMKII $\beta$  are abundantly expressed in [brain nerve cells](#) and have been extensively studied to show their important role in functions such as memory. However, the functions and conditions of CaMKII $\alpha$  and CaMKII $\beta$  in sleep control were unknown.

**CaMKII $\beta$  mutant (extended sleep)**



**CaMKII inhibitor (reduced sleep)**

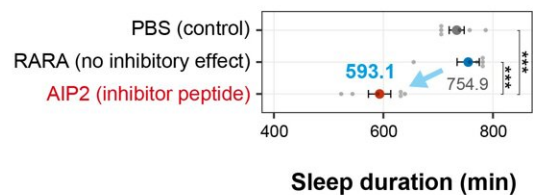


Fig.2 Bidirectional regulation of sleep by CaMKII The T287D mutant extends sleep duration, while mutations that do not act as phosphomimetics (T287A), and a mutation that causes loss of enzyme activity (K43R), indicating that the T287 phosphorylation and CaMKII enzyme activity is important for sleep regulation (left). Expression of a CaMKII inhibitor peptide resulted in a shortened sleep duration (right). Credit: Tone et al.

It is well-known that CaMKII $\beta$  has multiple phosphorylation sites and that its function is modulated according to their phosphorylation status. Therefore, this [study group](#) generated a series of CaMKII $\beta$  mutants. Toward this, mutated residues that mimic the phosphorylation state were introduced at each site, and the relationship between CaMKII $\beta$  phosphorylation state and [sleep regulation](#) was comprehensively investigated using mice. The researchers found that in specific phosphorylation state, CaMKII $\beta$  promoted the transition from wakefulness to sleep state. Furthermore, when an additional second or third similarly phosphorylated variant was introduced into CaMKII $\beta$  in this specific phosphorylated state, transition from the sleep state to wakefulness was inhibited.

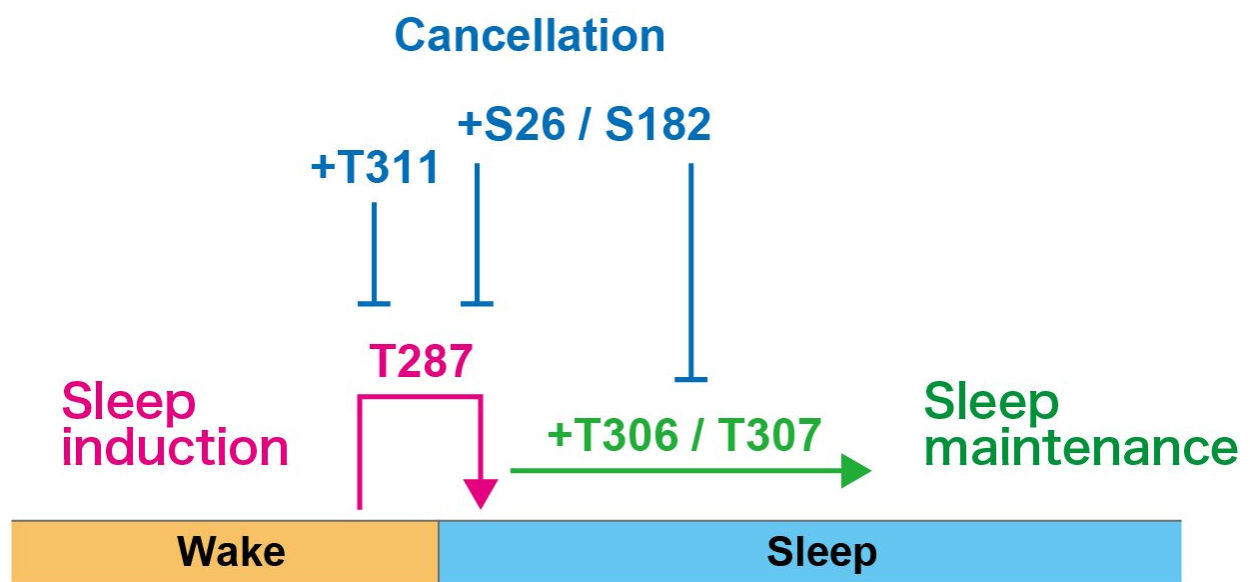


Fig.3 A schematic model of sleep regulation by CaMKII $\beta$  phosphorylation state. CaMKII $\beta$  acts on induction (T287), maintenance (T306, T307), and their cancellation (S26, S182, and T311) of sleep, depending on its own phosphorylation state. Credit: Tone et al.

In the present study, now published in *PLOS Biology*, they found that CaMKII $\beta$  functions at each step of a multi-step process that prolongs [sleep duration](#) (induction and maintenance of sleep) depending on its state of phosphorylation. Many existing hypnotics act on the steps involved in the induction of sleep. This study partly clarified the mechanism underlying sleep induction and maintenance and provides clues to consider more ideal sleep control methods aimed at maintaining sleep, which has been difficult so far.

This result was obtained as part of the JST Exploratory Research for Advanced Technology (ERATO) under the Ueda Biological Timing Project. In this project, JST developed an approach based on [systems biology](#) contributing to human understanding using sleep–wake rhythms as a model. The aim was to obtain information pertaining to biological timing that extends from molecules to individuals, in terms of human sleep–wake cycle.

**More information:** Daisuke Tone et al, Distinct phosphorylation states of mammalian CaMKII $\beta$  control the induction and maintenance of sleep, *PLOS Biology* (2022). [DOI: 10.1371/journal.pbio.3001813](https://doi.org/10.1371/journal.pbio.3001813)

Provided by Japan Science and Technology Agency

Citation: Elucidation of proteins controlling the sleep and awake transition (2022, October 26) retrieved 21 March 2023 from <https://phys.org/news/2022-10-elucidation-proteins-transition.html>

<p>This document is subject to copyright. Apart from any fair dealing for the purpose of private study or research, no part may be reproduced without the written permission. The content is provided for information purposes only.</p>
--