

SLEEP REGULATION

ppERK, sleep & fly

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Sleep is still a mysterious process. Its functions and regulatory mechanisms are unclear, but the fruit fly has emerged as a promising model system for uncovering its molecular underpinnings. Foltenyi *et al.* now show that a well-characterized pathway that involves epidermal growth factor receptor (EGFR) signalling regulates sleep in *Drosophila*.

The EGFR signalling pathway is similar in mammals and *Drosophila*. The transforming growth factor- α (TGF- α) homologues *Spitz*, *Gurken* and *Keren* (in *Drosophila*) can be processed by the transmembrane protein *Star* and the protease Rhomboid (*Rho*) and bind to EGFR, leading to activation of extracellular signal-regulated kinase (ERK).

The authors found that temporally increasing the expression of *star* and *rho* in fruit flies caused an increase in their sleep time, followed by a compensatory decrease. *Rho* and *Star* act upstream of EGFR and, thus, expressing a dominant-negative form of the receptor abolished their effects on sleep.

The authors used RNA interference (RNAi) to generate flies in

which *Rho* activity was reduced (*rho*^{DN} flies), first targeting the RNAi to all neurons and then to a restricted set of cells with the neural drivers *c767*, *50Y* and *c687*. They found that this dramatically reduced total sleep levels, with a higher number but shorter duration of sleep bouts, indicating that flies with low *Rho* levels still required sleep but could not maintain it. The authors next induced expression of secreted *Spitz* in the same restricted set of neurons. This increased sleep time by 1 hour, further confirming that cells that are driven by *c767*, *50Y* and *c687* are involved in sleep regulation.

On the molecular level, the authors measured ERK activation in flies that overexpressed both *rho* and *star*, and found that high levels of phosphorylated ERK correlated with an increase in sleep time courses. No ERK hyperphosphorylation occurred in flies that co-expressed *rho* and *star* with a dominant-negative form of EGFR, confirming that ERK activation occurs downstream of EGFR activation. The authors further found that the high level of phosphorylated ERK that resulted



from *rho* and *star* activation clustered to the tritocerebrum in the fly brain. The neural drivers *c767*, *50Y* and *c687* were localized within the pars intercerebralis, which projects to the tritocerebrum. The pars intercerebralis might be the fly equivalent of the mammalian hypothalamus, a region of the brain that is known to regulate arousal.

This study shows that the EGFR–ERK pathway regulates the maintenance of sleep in *Drosophila*. This signalling pathway is common to mammals, suggesting that it might also have a role in sleep regulation in other species.

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