In This Issue

EGF takes dual control of mRNA

EGF coordinates the nuclear and cytoplasmic activities of an RNA-binding protein to ensure a specific mRNA is translated at the right time and place in neurons, say Tsai et al.

mRNAs localize to specific regions within neurons, where they can be translated to quickly generate large amounts of a protein in the place where it is needed. Due to the large size of neurons, this is much more efficient than translating all mRNAs in a single place and then shipping out each protein to its site of action. But the movement and translation of these mRNAs must be tightly regulated, potentially by extracellular signals such as growth factors.

A SOLO performance in meiotic cohesion

Yan et al. describe how a Drosophila protein works with the cohesin complex to hold sister chromatids together during meiosis, ensuring their faithful segregation into sperm.

Just as in mitosis, the cohesin complex keeps sister chromatids together until they separate in the second of meiosis' two divisions. In addition, sister chromatid cohesion helps homologous chromosomes pair up and divide in the first meiotic division. But the function of cohesins in Drosophila meiosis is largely unknown because mutations in the proteins are lethal, and flies lack a homologue of the meiosis-specific cohesin Rec8.

Identifying the prime suspects in vesicle release

Neurotransmitters are secreted rapidly after stimulation because synaptic vesicles wait at the presynapse with their membrane fusion machinery already part-assembled, say Walter et al.

Synaptic vesicles dock at the plasma membrane and become “primed” for quick release in response to increased calcium levels. The release stage is controlled by SNARE proteins in both the vesicle and presynaptic membranes, which assemble into a complex that promotes membrane fusion. Whether the vesicle SNARE synaptobrevin only binds its target membrane partners at the fusion step or whether it binds earlier during vesicle priming is unclear.

Walter et al. mutated synaptobrevin’s SNARE-interaction domain and measured the protein’s ability to support vesicle priming and fusion in neuroendocrine cells. Mutations at the N-terminal end that destabilized synaptobrevin’s interaction with its SNARE partners lowered the number of primed vesicles. But the vesicles that remained were still secreted as quickly as they were in cells expressing wild-type synaptobrevin. In contrast, mutations in the C-terminal end of synaptobrevin’s binding motif slowed the speed with which vesicles were released.

The researchers think that the interaction domains of synaptobrevin and the other SNARE proteins come together at their N termini to prime vesicles for release and pause in this intermediate state before calcium causes them to zip their C termini into a fully assembled SNARE complex driving membrane fusion. Once they are primed, synaptic vesicles might be committed to release, so senior author Jakob Sørensen now wants to investigate how the priming step is regulated.

Tsai, N.-P., et al. 2010. J. Cell Biol. doi:10.1083/jcb.200910083.

Yan, R., et al. 2010. J. Cell Biol. doi:10.1083/jcb.200904040.