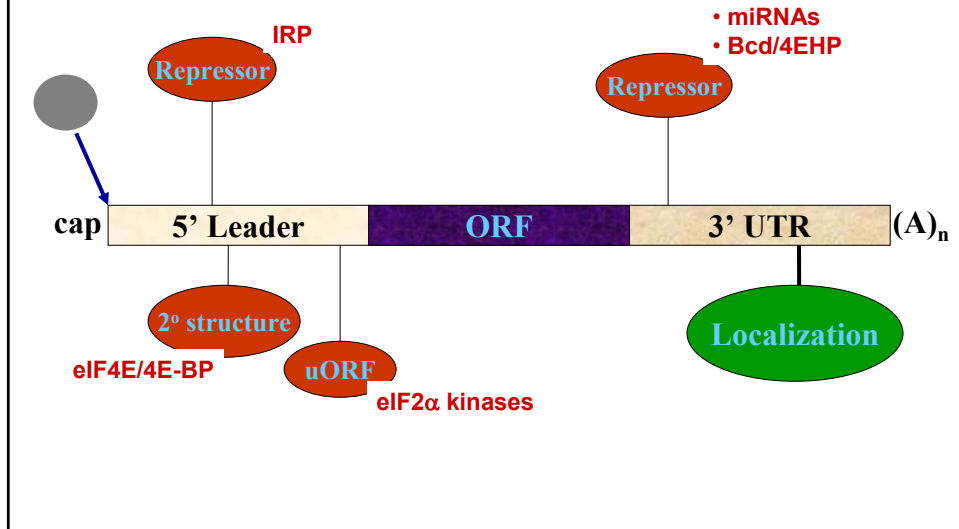
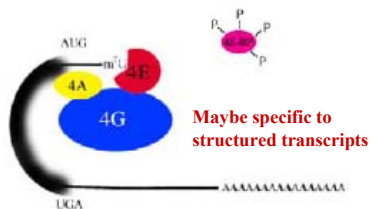


mRNA *cis* Elements Located in Non-Protein-Coding Regions

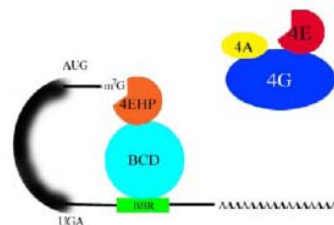


Transcript-independent versus specific competition at the 5' cap

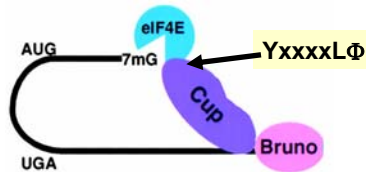
Transcript-independent inhibition by 4E-BPs



Transcript-specific inhibition by mRNA-bound bicoid



Other 3'-tethered repressors are 4E-BPs that complete with eIF-4G



Target mRNA	4E-BP	RNA-BP
<i>oskar</i>	Cup	Bruno
<i>nanos</i>	Cup	Smaug
cyclin B1*	Maskin	CPEB

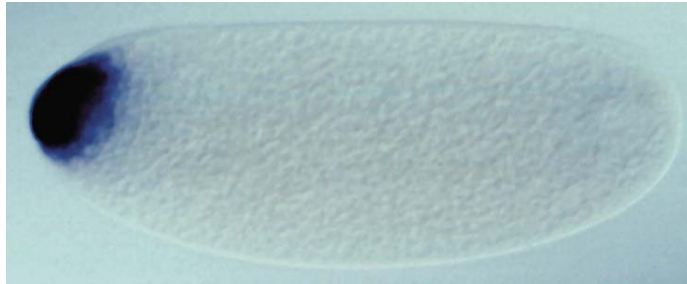
*Xenopus

Mechanisms of transcript-specific translational control

- Translational repressors (proteins, microRNAs) that bind to specific *cis* elements
- Interplay between structural characteristics of an mRNA (uORFs, secondary structure) and the activity of a general initiation factor (e.g. eIF2, eIF4E) can convert global to gene-specific regulation

Lecture 3

Mechanisms of Localized mRNA Translation in Cells



Reasons for mRNA Localization

- Targets protein to appropriate region of cell.
- Prevents expression elsewhere.
- Response to local requirements.
- Independent control in different regions.
- Localized synthesis necessary for assembly.
- More efficient transport
one mRNA vs. many proteins

Mechanisms of mRNA Localization

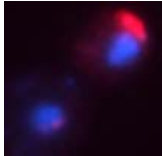
- Diffusion and localized entrapment.
- Localized degradation.
- Active transport on cytoskeletal motors.

Steps in mRNA localization

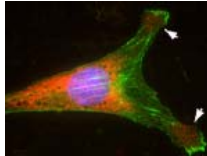
- Nuclear “priming” of transcripts.
- Translational silencing in cytosol.
- Specific association with motor proteins.
- Transport.
- Tethering at destination.
- Translational derepression.

Examples of mRNAs Actively Transported and Localized

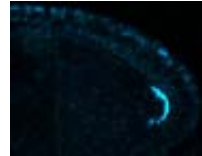
ASH1
Bud tip in dividing
S. cerevisiae



β-actin
Leading edge of
migrating fibroblast



oskar
Posterior of
drosophila oocyte



bicoid

ASH1

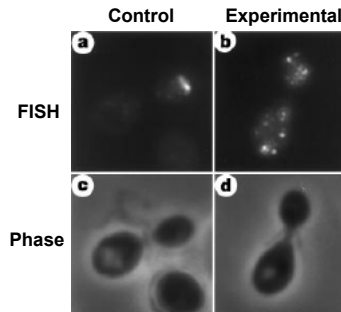


- Transcription factor
- Represses mating type switching in daughter cells.
- One of ~2 dozen mRNAs that localized to daughter cells
all seem to use the same mechanism

ASH1 mRNA travels along the actin cytoskeleton

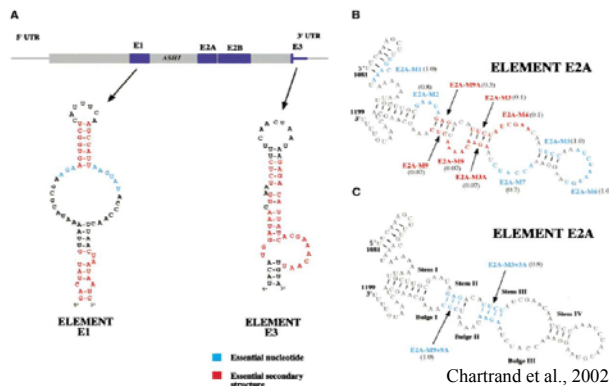
Delocalized by:

- *ACT1* mutants
- Actin assembly proteins (*SHE5*)
- *MYO4* mutants
- Actin depolymerization drugs



Takizawa et al., 1997

ASH1 cis-Elements

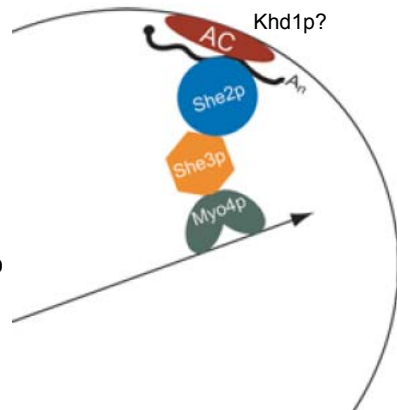


Chartrand et al., 2002

- Located in both the ORF and the 3'UTR
- Structure, not sequence, is important
- Inhibit translation as well as direct localization
- Each acts independently
 - Additive influence on *ASH1* localization
 - Each will localize a reporter gene individually

Model for *ASH1* mRNA localization

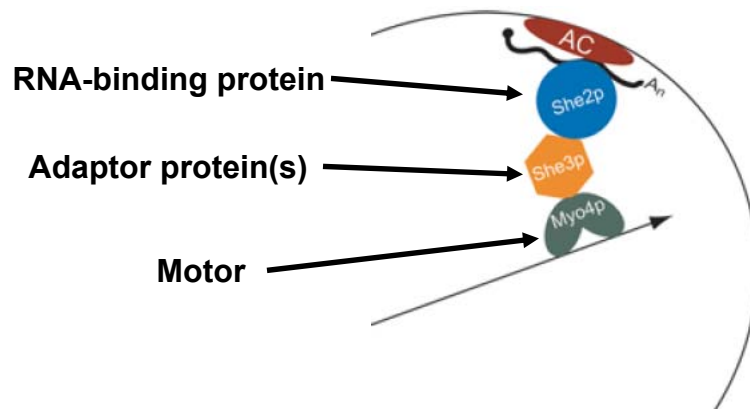
- **She2p**
RNA-binding protein
present in nucleus
- **Myo4p**
myosin motor protein
- **She3p**
interacts with She2p & Myo4p
required for Myo4p—She2p



- Multiple motors per transcript?
- Mechanism of anchoring?
- Activation of translation?

Gonsalvez et al., 2005

General Picture

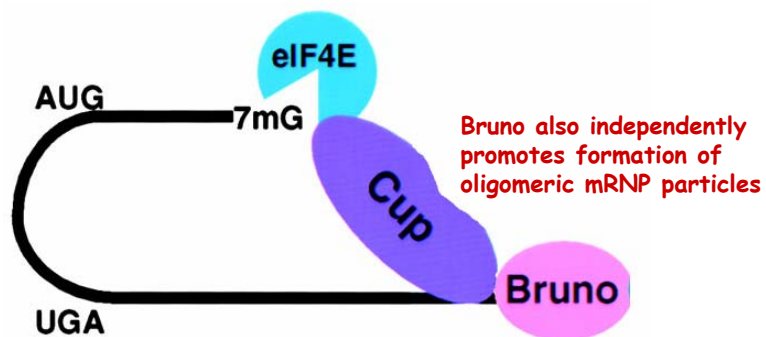


oskar



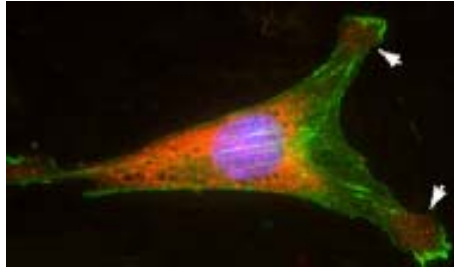
- Localized to posterior.
- Localization requires factors acquired in the nucleus.
- Cotransports with kinesin-HC (plus-end-directed MT motor).
- Not transported in *khc*⁻ mutants.
- **Staufen**
 - highly conserved dsRNA binding protein
 - required and colocalizes

Inappropriate *oskar* translation is inhibited by tethering eIF4E



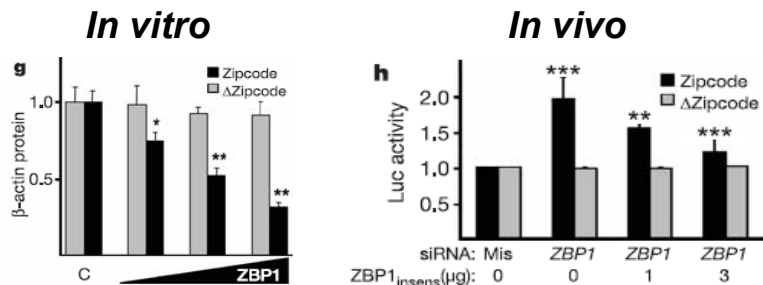
How is it regulated once the posterior site is reached?

β -Actin mRNA in Fibroblasts

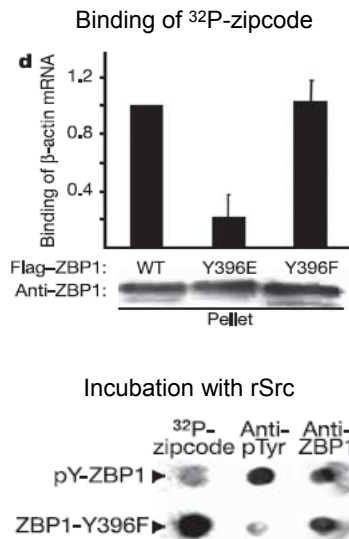


- Localizes to the leading edge of migrating cells.
- Mediated by two 54-nt elements in 3'UTR ("zipcodes").
- ZBP1 binds to the zipcodes and is required for transport.
- ZBP1 assembles onto the 3'UTR and travels with the *β -actin* mRNA.
- Active myosin and intact actin filaments are required.
- ZBP1 also inhibits translation of *β -actin* mRNA.

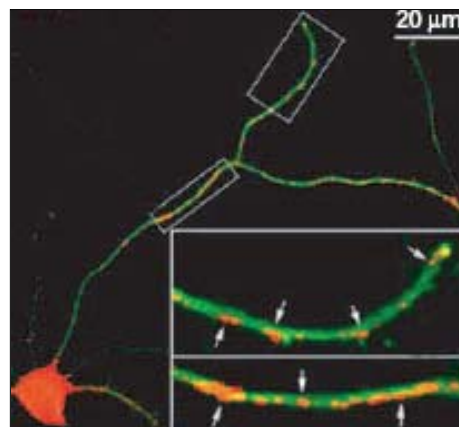
ZBP1 inhibits translation in a zipcode dependent manner



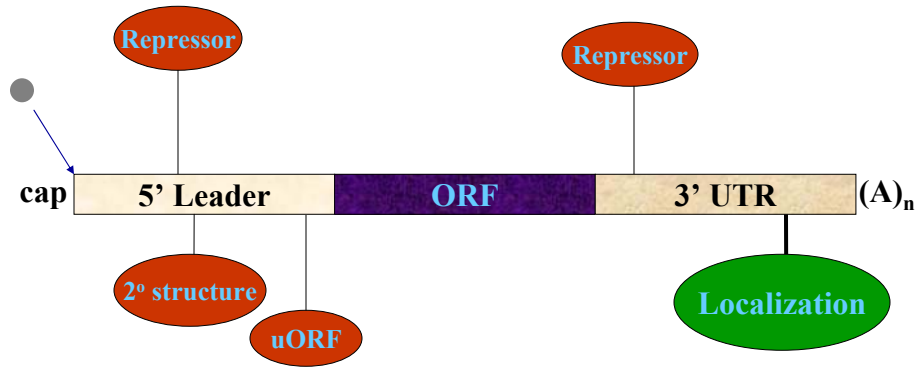
Phosphorylation of Tyr396 by Src kinase blocks interaction of ZBP1 with zipcode RNA



ZBP1 is also involved in β -actin mRNA movement in dendrites, but in contrast to fibroblasts, transport is on microtubules using kinesin motors.



mRNA *cis* Elements Located in Non-Protein-Coding Regions



Most of these regulatory systems do not involve simple one-protein/one-RNA interactions