

She2p is a novel RNA-binding protein that recruits the Myo4p–She3p complex to *ASH1* mRNA

Roy M. Long¹, Wei Gu², Ellen Lorimer,
Robert H. Singer² and Pascal Chartrand²

Medical College of Wisconsin, Department of Microbiology and Molecular Genetics, 8701 Watertown Plank Road, Milwaukee, WI 53226-0509 and ²Albert Einstein College of Medicine, Department of Anatomy and Structural Biology, 1300 Morris Park Avenue, Bronx, NY 10461, USA

¹Corresponding author
e-mail: rlong@mcw.edu

In *Saccharomyces cerevisiae*, Ash1p is a specific repressor of transcription that localizes exclusively to daughter cell nuclei through the asymmetric localization of *ASH1* mRNA. This localization requires four *cis*-acting localization elements located in the *ASH1* mRNA, five *trans*-acting factors, one of which is a myosin, and the actin cytoskeleton. The RNA-binding proteins that interact with these *cis*-elements remained to be identified. Starting with the 3' most localization element of *ASH1* mRNA in the three-hybrid assay, element E3, we isolated a clone corresponding to the C-terminus of She3p. We also found that She3p and She2p interact, and this interaction is essential for the binding of She3p with element E3 *in vivo*. Moreover, She2p was observed to bind the E3 RNA directly *in vitro* and each of the *ASH1* *cis*-acting localization elements requires She2p for their localization function. By tethering a She3p–MS2 fusion protein to a reporter RNA containing MS2 binding sites, we observed that She2p is dispensable for She3p–MS2-dependent RNA localization.

Keywords: *ASH1*/mRNA localization/RNA-binding proteins/*Saccharomyces cerevisiae*/three-hybrid assay

Introduction

The process of cellular differentiation during development of multicellular organisms requires the asymmetric sorting of cell fate determinants between sister cells. One mechanism used for the asymmetric segregation of cell fate determinants is to localize specifically the mRNAs encoding the determinants to one of the sister cells. The currently emerging mechanism for the localization of RNA involves the presence of specific *cis*-acting localization elements within the localized mRNA. The *cis*-acting elements are recognized by RNA-binding proteins that physically interact with molecular motors that transport the ribonucleoprotein complex along cytoskeletal filaments to the site of localization (Nasmyth and Jansen, 1997; Oleynikov and Singer, 1998; Schnorrer *et al.*, 2000).

The asymmetric sorting of Ash1p in the yeast *Saccharomyces cerevisiae* is a model system for studying the asymmetric segregation of cell fate determinants

through RNA localization. Ash1p is a cell-type specific transcriptional repressor of the HO endonuclease (Bobola *et al.*, 1996; Sil and Herskowitz, 1996). At the end of mitosis, Ash1p is found in the nucleus of daughter cells where it represses HO transcription and ultimately prevents mating-type switching in these cells (Bobola *et al.*, 1996; Sil and Herskowitz, 1996). This transcriptional regulation of HO expression restricts mating-type switching to mother cells (Bobola *et al.*, 1996; Sil and Herskowitz, 1996). The asymmetric distribution of Ash1p results from the localization of *ASH1* mRNA to the distal tip of daughter cells during anaphase (Long *et al.*, 1997; Takizawa *et al.*, 1997).

ASH1 mRNA contains four *cis*-acting localization elements: E1, E2A, E2B and E3 (Chartrand *et al.*, 1999; Gonzalez *et al.*, 1999). Each of these elements is sufficient to localize a heterologous reporter mRNA to daughter cells. The E3 element is located predominantly in the 3' untranslated region (3'-UTR) whereas the remaining three elements are found in the *ASH1* open reading frame (ORF). The E1 and E3 elements are predicted to fold into structures containing stem-loops. Mutations predicted to disrupt the stem-loop structure of either element prevent RNA localization activity, while compensatory mutations that re-establish the secondary structure restore RNA localization activity (Chartrand *et al.*, 1999; Gonzalez *et al.*, 1999). For the E3 element, specific RNA sequences have not been demonstrated to be important for RNA localization activity; rather, RNA localization activity for E3 is strictly dependent on secondary structure (Chartrand *et al.*, 1999; Gonzalez *et al.*, 1999). In contrast, in E1, specific RNA sequences important for RNA localization activity have been identified (Chartrand *et al.*, 1999).

Five genes, *SHE1–5*, originally found to be essential for proper sorting of Ash1p, are also required for *ASH1* mRNA localization (Jansen *et al.*, 1996; Long *et al.*, 1997; Takizawa *et al.*, 1997). Several observations indicate that actomyosin-based transport is important for *ASH1* mRNA localization (Long *et al.*, 1997; Takizawa *et al.*, 1997). A functional actin cytoskeleton was found to be essential for *ASH1* mRNA localization and *SHE5/BNL1* is required for proper organization of the actin cytoskeleton (Kohnno *et al.*, 1996; Evangelista *et al.*, 1997). *SHE4* is also hypothesized to be required for proper organization of the actin cytoskeleton (Wendland *et al.*, 1996). Furthermore, *SHE1/MYO4* encodes a type V myosin, which co-localizes with *ASH1* mRNA at the tip of daughter cells (Haarer *et al.*, 1994; Münchow *et al.*, 1999; Takizawa and Vale, 2000). Myo4p was also found to be associated with *ASH1* mRNA in co-immunoprecipitation experiments (Münchow *et al.*, 1999; Takizawa and Vale, 2000). A live cell assay has shown particles containing Myo4p and *ASH1* mRNA rapidly moving from mother cells to daughter cells,

suggesting a direct role for Myo4p in *ASH1* mRNA transport (Bertrand *et al.*, 1998; Beach *et al.*, 1999).

While the role of myosin in the localization of *ASH1* mRNA has apparently been determined, a direct interaction between *ASH1* mRNA and Myo4p has not been established. On the contrary, new studies suggest that the association of Myo4p with *ASH1* mRNA requires She2p and She3p (Münchow *et al.*, 1999; Takizawa and Vale, 2000). By two-hybrid analysis, Myo4p and She3p have been shown to interact (Münchow *et al.*, 1999). *ASH1* mRNA can be co-immunoprecipitated in a complex containing Myo4p, She2p and She3p. Co-immunoprecipitation of *ASH1* mRNA with Myo4p and She3p is dependent on She2p (Münchow *et al.*, 1999; Takizawa and Vale, 2000); however, these experiments provide no definite function for either She2p or She3p and do not address what additional unknown proteins may be co-immunoprecipitating with the complex.

Using molecular genetics and biochemical assays, we demonstrate directly that She2p is an RNA-binding protein that interacts specifically with individual *ASH1* mRNA localization elements and functions to recruit the Myo4p–She3p complex to *ASH1* mRNA.

Results

Interaction of She3p with an *ASH1* mRNA localization element

Our approach was to identify specific RNA-binding proteins for *ASH1* *cis*-acting localization elements and to use these RNA-binding proteins to characterize protein–protein interactions required for *ASH1* mRNA localization. We initiated an *in vivo* approach for the identification of RNA-binding proteins using the three-hybrid assay (SenGupta *et al.*, 1996). In this system, formation of an RNA–protein complex in yeast results in transcriptional activation of the reporter genes *HIS3* and *lacZ*.

Plasmid pRL80, expressing a fusion RNA containing two copies of the MS2 coat protein stem–loop motif and one copy of the 127 nucleotide (nt) *ASH1* 3′-UTR localization element (E3), was transformed into the three-hybrid host yeast strain YBZ-1. A yeast cDNA library fused to the Gal4 transcriptional activation domain (Gal4p-AD) was screened for proteins that could interact with the *ASH1* E3 localization element. From 500 000 total yeast transformants, we isolated one clone corresponding to the C-terminus (amino acids 236–425) of She3p. *SHE3* encodes a protein of 47 kDa with no significant homology to other known proteins (Jansen *et al.*, 1996). In subsequent three-hybrid screens, She3p was repeatedly isolated, but none of these clones contained the entire She3p ORF.

The binding specificity of the *SHE3* C-terminal clone was investigated by three-hybrid analysis using mutants in the E3 element (M9, M13, M14) that are predicted to disrupt the secondary structure of element E3 and are defective for RNA localization (Chartrand *et al.*, 1999). We observed in the three-hybrid assay that She3p C-terminal-dependent β -galactosidase expression was lower in mutants M9, M13 and M14 compared with the wild-type E3 element (Figure 1A). In contrast, an E3

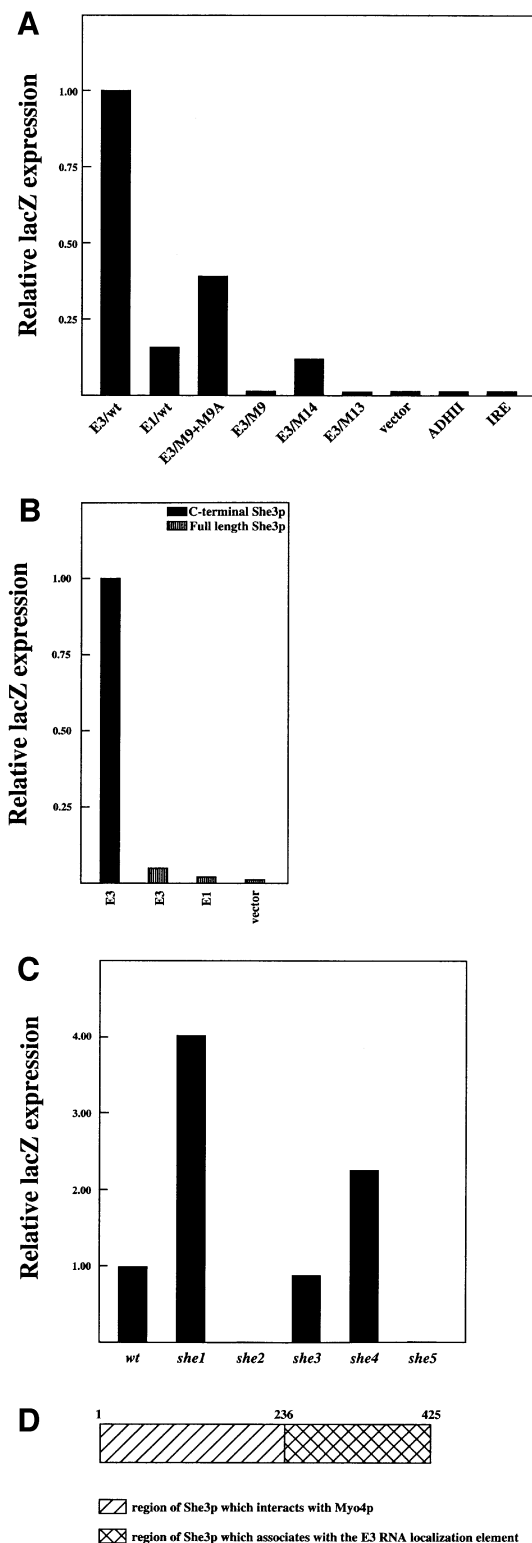
double mutant (M9 + M9A), predicted to restore the wild-type stem–loop structure and capable of RNA localization, resulted in higher She3p C-terminal-dependent β -galactosidase expression (Figure 1A) compared with the M9, M13 and M14 mutations (Chartrand *et al.*, 1999). When other yeast 3′-UTRs (ADHII) or double-stranded RNAs, like the iron-responsive element (IRE) (Klausner *et al.*, 1993; Melefors and Hentze, 1993), were investigated for She3p C-terminal-dependent β -galactosidase expression, we observed expression levels similar to the vector control (Figure 1A). These results indicate that the She3p interaction is specific for the E3 localization element. These data suggest that the interaction of the C-terminus of She3p with element E3 is apparently dependent on a specific secondary structure of the E3 element. The sum of this work demonstrates a correlation between mRNA localization activity of E3 and the interaction of She3p with E3.

The observation that She3p is required for *ASH1* mRNA localization suggests that She3p should interact with each of the *ASH1* *cis*-acting localization elements. We tested the ability of the C-terminus of She3p to interact with the E1 element by three-hybrid analysis. We observed that She3p C-terminal-dependent β -galactosidase expression was higher with element E3 than element E1 (Figure 1A). The different levels of β -galactosidase expression between E1 and E3 could result from different expression levels of the three-hybrid RNA substrates. However, data obtained previously with the three-hybrid assay indicate that the expression levels of these two fusion RNAs are comparable (data not shown). Therefore, we conclude that the association between the C-terminus of She3p and elements E1 and E3 somehow differs.

The association between the C-terminus of She3p and the various *cis*-acting elements may differ from full-length She3p. Therefore, we investigated the interaction of the full-length She3p with the elements E1 and E3 (Figure 1B). Full-length She3p was cloned into the appropriate three-hybrid vector and the level of β -galactosidase expression was determined for full-length She3p with either E1 or E3 fusion RNAs. We observed only a slight increase in β -galactosidase expression over the vector control (Figure 1B). Furthermore, the C-terminus of She3p displayed higher β -galactosidase expression levels with E3 compared with full-length She3p (Figure 1B). These results suggested that the N-terminus of She3p negatively affected the association between She3p and the *cis*-acting elements as assayed by the three-hybrid system.

One possibility could be that full-length She3p was bound to a cytoplasmic factor preventing She3p from entering the nucleus in the three-hybrid assay. Since She3p interacts with Myo4p in the two-hybrid assay (Münchow *et al.*, 1999), this interaction could directly or indirectly affect the ability of full-length She3p to associate with the three-hybrid RNA. We tested the interaction of full-length She3p with element E3 in a *myo4* strain and observed a 4-fold increase in β -galactosidase expression levels compared with the *MYO4* strain (Figure 1C). This result indicates that Myo4p prevented full-length She3p from associating with the *cis*-acting localization elements in the three-hybrid system. Presumably, Myo4p associates with cytoplasmic actin and full-length She3p, through a domain of She3p between amino acids 1 and 235, making She3p

unavailable for associating with nuclear RNA in the three-hybrid assay. In summary, She3p appears to contain at least two important domains: one responsible for interaction with Myo4p and a second independent domain required for an association with *ASH1* mRNA *cis*-acting localization sequences (Figure 1D).



The *in vivo* interaction of She3p with the E3 localization sequence is dependent on She2p

The *in vivo* association between *ASH1* mRNA, Myo4p and She3p is dependent on She2p (Münchow *et al.*, 1999; Takizawa and Vale, 2000). We sought to determine whether the *in vivo* association of *ASH1* mRNA and the C-terminus of She3p was dependent on She2p or any of the other She proteins. Deletions of *SHE1–5* were created in strain YBZ-1, and C-terminal She3p–E3-dependent β -galactosidase expression levels were determined (Figure 2). We observed that deletion of *MYO4* or *SHE3* had no significant effect on β -galactosidase expression, and deletion of *SHE4* or *SHE5* resulted in a modest reduction in β -galactosidase activity (Figure 2). In contrast, deletion of *SHE2* resulted in a dramatic reduction in β -galactosidase expression, indicating that the *in vivo* association between She3p and E3 requires She2p. By western and northern blotting for the three-hybrid

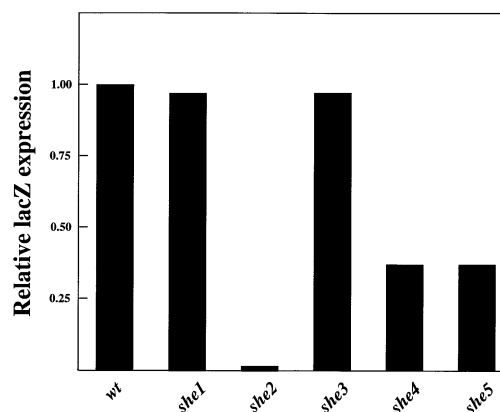


Fig. 2. She3p C-terminal-dependent lacZ expression assayed by three-hybrid analysis is dependent on She2p. Yeast strain YBZ-1 deleted of *myo4*, *she2*, *she3*, *she4* and *she5*. These strains were subsequently transformed with a plasmid expressing the C-terminus She3p fused to Gal4p-AD and MS2–E3 fusion RNA. The transformants were grown in liquid culture and processed for β -galactosidase assays. β -galactosidase expression levels were normalized to the wild-type strain.

Fig. 1. By three-hybrid analysis, the C-terminal section of She3p preferentially interacts with the E3 *ASH1* RNA localization element. (A) Strain YBZ-1 transformed with a plasmid expressing Gal4p-AD fused to the C-terminus of She3p along with a plasmid expressing one of the indicated MS2 fusion RNAs. Liquid cultures were grown for each strain, and the cells harvested by centrifugation. The cells were permeabilized and β -galactosidase expression levels were determined. β -galactosidase expression levels were normalized to cells expressing the MS2–E3 fusion RNA. (B) Strain YBZ-1 transformed with either a plasmid expressing the C-terminus of She3p fused to Gal4p-AD or full-length She3p fused to Gal4p-AD along with plasmids expressing the indicated MS2 fusion RNAs. Liquid cultures were grown for each strain and used for β -galactosidase assays. β -galactosidase expression levels were normalized to cells expressing the She3p C-terminus fused to Gal4p-AD and the MS2–E3 fusion RNA. (C) Yeast strain YBZ-1 deleted of *myo4*, *she2*, *she3*, *she4* or *she5*. The strains were subsequently transformed with a plasmid expressing full-length She3p fused to Gal4p-AD along with a plasmid expressing the MS2–E3 fusion RNA. Transformants were grown in liquid culture and used for β -galactosidase assays. β -galactosidase levels were normalized to the wild-type strain. (D) Schematic representation of She3p. Amino acids 1–236 of She3p contain a domain required for the association with Myo4p, while amino acids 236–425 contain a domain required for the *in vivo* association with the E3 *ASH1* RNA localization element.

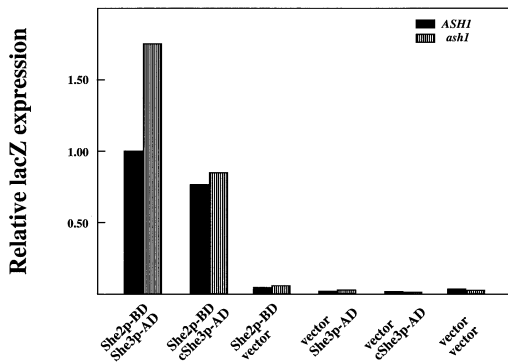


Fig. 3. By the two-hybrid assay, She2p and She3p interact. Yeast strains PJ69-4a (*ASH1*) and YLM445 (*ash1*) were transformed in combinations with the following plasmids: She2p fused to Gal4p-BD (She2p-BD), full-length She3p fused to Gal4p-AD (She3p-AD), the C-terminus of She3p fused to Gal4p-AD (cShe3p-AD), the vector corresponding to Gal4p-BD (pGBDU-c2) or the vector corresponding to Gal4p-AD (pACT2). Transformants were grown in liquid culture and processed for β -galactosidase assays. β -galactosidase expression levels were normalized to strain PJ69-4a transformed with the plasmid expressing She2p fused to Gal4p-BD and the plasmid expressing full-length She3p fused to Gal4p-AD.

components we eliminated the possible trivial explanation that the abundance of the three-hybrid components is reduced in the *she2* strain compared with the wild-type control (data not shown). These results suggest that She2p has a direct role in recruiting She3p to *ASH1* mRNA localization sequences.

Association of She2p and She3p

Since the *in vivo* association between She3p and the E3 localization element was dependent on She2p, we reasoned that She2p and She3p should also associate *in vivo*. We investigated the association between She2p and She3p by two-hybrid analysis (Fields and Song, 1989). She2p was fused to the Gal4 DNA-binding domain (Gal4p-BD). This She2p bait was tested against the entire ORF for She3p or the C-terminus of She3p when either was fused to Gal4p-AD. We only observed activation of *lacZ* expression when yeast was simultaneously transformed with She2p-Gal4p-AD and either She3p-Gal4p-BD construct (Figure 3). From these results we conclude that She2p and She3p interact either directly or indirectly, and the domain of She3p responsible for the interaction with She2p resides between amino acids 236 and 425 of She3p. This is the same region of She3p responsible for the *in vivo* association with the E3 localization element.

The interaction between She2p and She3p could occur indirectly through *ASH1* mRNA tethering the two proteins together. We deleted *ASH1* from the two-hybrid strain and assayed She2p-She3p-dependent β -galactosidase expression (Figure 3). In the absence of *ASH1* mRNA we measured a modest increase in β -galactosidase expression, providing evidence for a direct association between She2p and She3p.

She2p RNA-binding activity

The region of She3p required for the association with *ASH1* RNA localization elements was identical to that required for the association with She2p; we reasoned that

She2p could be the RNA-binding protein that recruits She3p to *ASH1* RNA localization elements. We tested She2p RNA-binding activity directly by expressing and purifying She2p from *Escherichia coli* as a glutathione *S*-transferase (GST) fusion protein. Purified GST-She2p was tested for RNA-binding activity by a UV cross-linking assay. When 32 P-labeled E3 was incubated with recombinant GST-She2p followed by UV cross-linking, SDS-PAGE and autoradiography, we observed a product corresponding closely to the molecular weight of the GST-She2p fusion protein (49.6 kDa) (Figure 4A). We did not observe this product in the presence of 32 P-labeled pGEM-4Z RNA (Figure 4A). Furthermore, the GST-She2p UV cross-linked product was specifically competed with 1000 \times excess unlabeled E3 and not with 1000 \times excess unlabeled pGEM-4Z RNA sequences (Figure 4B). We have further observed that 5000 \times excess pGEM-4Z RNA will not compete the She2p-E3 UV cross-linked product (data not shown).

The specificity of the She2p-E3 interaction was further examined using mutations in the E3 localization element. Mutations in the E3 element (M9, M13 and M14) previously shown to be defective in RNA localization did not compete the She2p-E3 UV cross-linked product, while a mutation in E3 (M9 + M9A) that supports RNA localization did compete the She2p-E3 UV cross-linked product (Figure 4C). These results demonstrate that She2p is capable of directly and specifically binding the E3 localization element.

We investigated by UV cross-linking whether She2p could interact with each of the *cis*-acting localization elements using recombinant GST-She2p, 32 P-labeled E3 and excess unlabeled RNA for each element. We observed that 1000 \times excess unlabeled RNA corresponding to either *ASH1* element E1 or E3 effectively competed the She2p-E3 UV cross-linked product (Figure 4D). However, a 1000-fold excess of elements E2A and E2B did not compete as effectively as E1 or E3 (Figure 4D). From these results we conclude that, *in vitro*, She2p is able to bind each of the *cis*-acting localization elements, but apparently with different affinities.

She2p is essential for the localization function of the four *ASH1* mRNA *cis*-acting elements

Analogous to She3p, She2p is required for the localization of full-length *ASH1* mRNA (Long *et al.*, 1997; Takizawa *et al.*, 1997), but the requirement for She2p by each of the *cis*-acting localization elements remained to be determined. The She2p RNA-binding data predict that RNA localization directed by each of the *cis*-acting elements should be dependent on She2p. To test this prediction, a wild-type strain and a *she2* strain were transformed with each of the *lacZ* mRNA reporter plasmids: *lacZ-ASH1-E1*, *lacZ-ASH1-E2A*, *lacZ-ASH1-E2B* and *lacZ-ASH1-E3*. The intracellular distribution of the *lacZ* RNA expressed from each of these plasmids was determined by FISH. For all of these reporter constructs we observed that the *lacZ* mRNA was localized to daughter cells in the wild-type strain and delocalized between mother and daughter cells in the *she2* strain (Figure 5). From these results we conclude that the activity of each *ASH1* *cis*-acting localization element requires She2p.

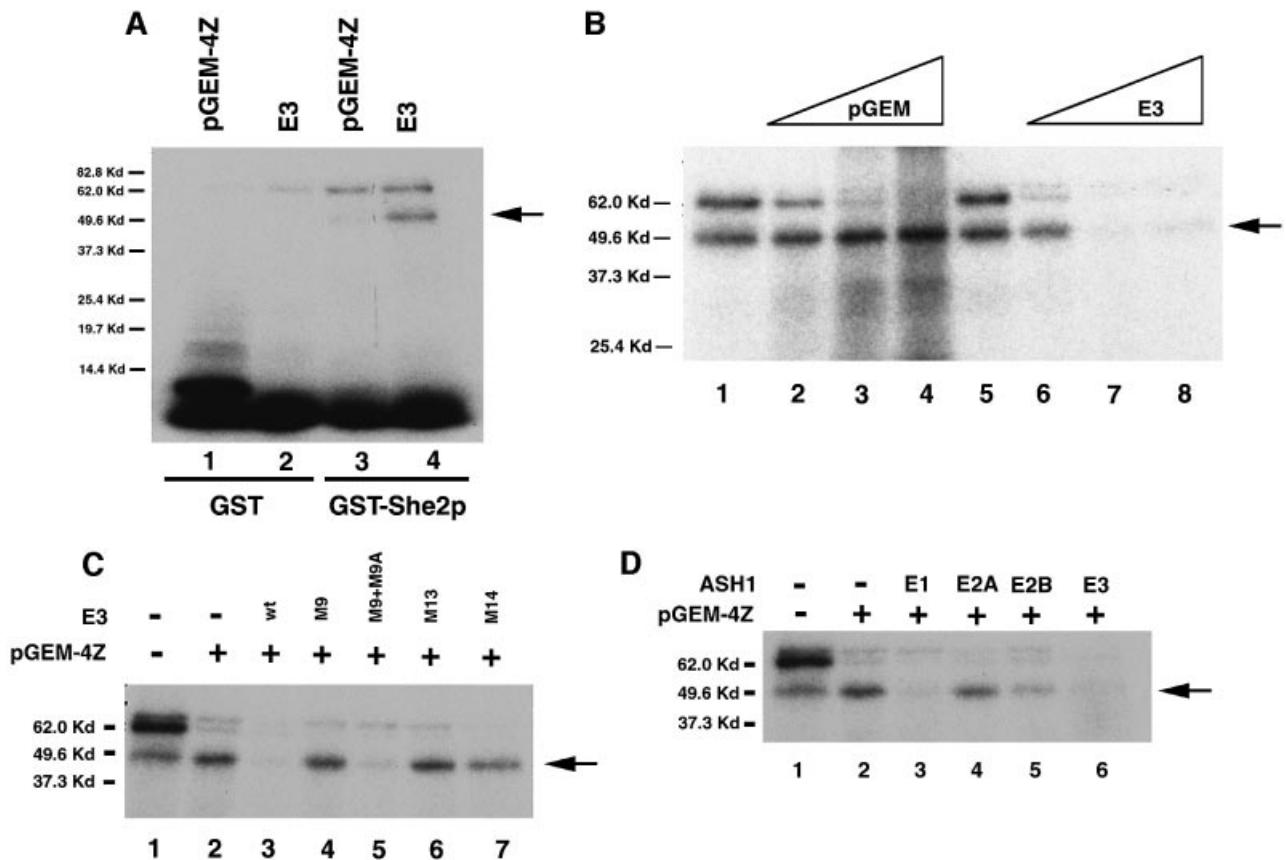


Fig. 4. She2p RNA-binding activity as assayed by a UV cross-linking assay. **(A)** One nanogram of ^{32}P -labeled E3 RNA (lanes 2 and 4) or pGEM-4Z RNA (lanes 1 and 3) was incubated with 1.0 μg of purified GST-She2p (lanes 3 and 4) or GST (lanes 1 and 2) corresponding to an equivalent volume of GST-She2p used for UV cross-linking. The GST-She2p-E3 complex migrates at 49.6 kDa. The arrow indicates the position of the GST-She2p-E3 complex. **(B)** Specific cross-linking of GST-She2p to E3. UV cross-linking was performed with GST-She2p and ^{32}P -labeled E3 in the absence (lanes 1 and 5) or presence of 100 \times (lane 2), 500 \times (lane 3) and 1000 \times (lane 4) pGEM-4Z RNA, as well as 100 \times (lane 6), 500 \times (lane 7) and 1000 \times (lane 8) E3 RNA. **(C)** Competition of GST-She2p-E3 UV cross-linked product with mutant E3 elements. GST-She2p was incubated with ^{32}P -labeled E3 in the absence of pGEM and E3 RNA (lane 1) or 1000 \times pGEM RNA (lane 2), 1000 \times pGEM/1000 \times E3 (lane 3), 1000 \times pGEM/1000 \times E3/M9 (lane 4), 1000 \times pGEM/1000 \times E3/M9 + M9A (lane 5), 1000 \times pGEM/1000 \times E3/M13 (lane 6) or 1000 \times pGEM/1000 \times E3/M14 (lane 7). **(D)** Competition of GST-She2p-E3 UV cross-linked product with *ASH1* cis-acting localization elements. GST-She2p was incubated with ^{32}P -labeled E3 in the absence of pGEM and E3 RNA (lane 1), 1000 \times pGEM RNA (lane 2), 1000 \times pGEM/1000 \times E1 (lane 3), 1000 \times pGEM/1000 \times E2A (lane 4), 1000 \times pGEM/1000 \times E2B (lane 5) or 1000 \times pGEM/1000 \times E3 (lane 6).

***She2p* functions to recruit *Myo4p*-*She3p* to *ASH1* RNA localization elements**

Our work, in addition to the work of others, implies that She2p's sole function in *ASH1* mRNA localization is to interface the Myo4p-She3p complex with *ASH1* mRNA. If this is the only function of She2p in Myo4p-She3p-dependent RNA localization, She2p should be dispensable for RNA localization if She3p is given a high affinity RNA-binding domain. We tested this hypothesis by fusing She3p to the MS2 phage RNA-binding coat protein (Fouts *et al.*, 1997) and monitoring the localization of a lacZ reporter RNA containing six MS2 binding sites in its 3'-UTR (Bertrand *et al.*, 1998). In wild-type cells, we observed that lacZ mRNA was localized to the tip of daughter cells dependent on the presence of MS2 binding sites in the reporter mRNA as well as the expression of She3p-MS2 (Figure 6, compare A-C and D-F with G-I). Furthermore, localization of lacZ-MS2 by She3p-MS2 was dependent on Myo4p (Figure 6M-O). However, we observed in *she2* cells that She3p-MS2 was still able to localize lacZ-MS2 RNA to daughter cells (Figure 6J-L).

These results imply that the sole function of She2p in She3p-dependent RNA localization is to recruit Myo4p-She3p to *cis*-acting localization elements.

Discussion

We have identified and characterized a new RNA-binding protein required for *ASH1* mRNA localization in the yeast *S.cerevisiae*. Previously, it has been shown that Myo4p, She3p and *ASH1* mRNA are associated in a complex containing She2p (Bertrand *et al.*, 1998; Münchow *et al.*, 1999; Takizawa and Vale, 2000); however, a definitive role for She2p and She3p in this complex remained to be determined. Here we show for the first time that She2p can directly and specifically bind *ASH1* cis-acting localization elements. These results strongly suggest that She2p is the RNA-binding protein that interfaces *ASH1* mRNA to She3p and the transport motor, Myo4p.

Using a segment of *ASH1* mRNA (E3) shown to be sufficient for localization of mRNA to the bud in a three-hybrid assay, we found that She3p, a cytoplasmic protein

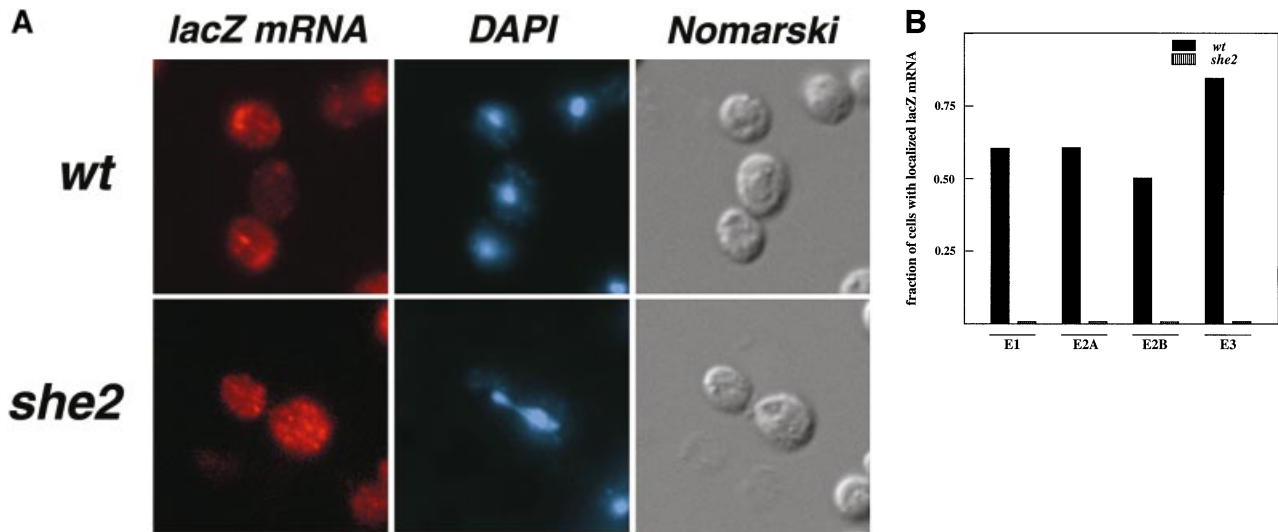


Fig. 5. RNA localization activity for each of the *cis*-acting localization elements is dependent on She2p. (A) Wild-type (wt) and *she2* yeast strains were transformed with a plasmid containing the galactose-inducible *lacZ-E3* cassette, and the cells were processed for fluorescent *in situ* hybridization (FISH). (B) Fraction of budding cells with localized *lacZ* mRNA for each of the *cis*-acting localization elements in wild-type and *she2* cells. Cells expressing each of the reporter constructs (*lacZ-E1*, *lacZ-E2A*, *lacZ-E2B* and *lacZ-E3*) were grown in galactose-containing media and processed for *in situ* hybridization. *LacZ* RNA was categorized as localized or delocalized from cells displaying FISH signal.

required for *ASH1* mRNA localization, associated with this RNA (Jansen *et al.*, 1996; Long *et al.*, 1997; Takizawa *et al.*, 1997). In this *in vivo* assay, we observed that She3p–E3-dependent β -galactosidase expression was dependent on a functional E3 element, since mutations in E3 (M9, M13 and M14) defective for RNA localization displayed significantly lower levels of β -galactosidase expression. Interestingly, the She3p clone isolated from the three-hybrid assay contained only a C-terminal portion of She3p. Full-length She3p was not isolated in the three-hybrid assay, and when full-length She3p was tested in the assay, β -galactosidase expression was greatly reduced compared with that observed for the C-terminus of She3p. We and others (R.-P.Jansen, personal communication) have found that full-length She3p, via its N-terminal region, is retained in the cytoplasm through protein–protein interactions with Myo4p. Since the three-hybrid analysis requires that the RNA–protein complex be imported into the nucleus for transcriptional activation to occur, anchoring of full-length She3p in the cytoplasm prevents activation of reporter gene expression.

However, the three-hybrid results do not prove that She3p directly binds the E3 localization element. Using recombinant She3p in a mobility shift assay, binding of She3p with element E3 RNA was not comparable in affinity to She2p, suggesting that She3p has low, if any, intrinsic RNA-binding activity. We favor the model whereby She3p interacts *in vivo* with an RNA-binding protein, She2p, specific for the *ASH1* mRNA localization sequences.

Like She3p, She2p is essential for the localization of *ASH1* mRNA (Long *et al.*, 1997; Takizawa *et al.*, 1997). She2p is also known to be essential for the association of Myo4p and She3p with *ASH1* mRNA (Münchow *et al.*, 1999; Takizawa and Vale, 2000). Moreover, She2p was found to be essential for the proper formation of a green fluorescent protein (GFP)–*ASH1* mRNA localization

particle, the ‘locasome’ (Bertrand *et al.*, 1998). Our results also show that the three-hybrid interaction of She3p with element E3 is dependent on She2p, and that She2p interacts with a domain in the C-terminus of She3p in the absence of *ASH1* mRNA. The possibility remains that She2p and She3p could be tethered together *in vivo* by an unidentified mRNA. These data indicate that She2p could possess RNA-binding activity and *in vivo* bind *ASH1* mRNA localization elements and recruit the Myo4p–She3p complex to *ASH1* mRNA.

Using recombinant She2 protein in a UV cross-linking assay, we observed specific binding of She2p with element E3 RNA, indicating that She2p is an RNA-binding protein. She2p’s binding to element E3 is affected by mutations in the secondary structure of this element. Furthermore, elements E1 and E3 successfully compete the element E3 for binding to She2p, suggesting that She2p recognizes diverse RNA secondary structures.

Furthermore, we have shown that each of the *cis*-acting elements requires She2p for RNA localization activity. This result supports the conclusion that an identical RNA localization complex assembles on each of the *cis*-acting localization elements. This conclusion also implies that each of the *cis*-acting localization elements performs redundant functions and the presence of multiple *cis*-acting elements in *ASH1* mRNA simply serves to increase the efficiency of *ASH1* RNA localization to daughter cells.

Model for *ASH1* mRNA localization

Combining this study with previous work, we propose the following model for *ASH1* mRNA localization (Figure 7). *ASH1* mRNA is identified in the nucleus by Loc1p, a nuclear RNA-binding protein required for *ASH1* mRNA localization (R.M.Long, W.Gu, G.Gonsalvez, R.H.Singer and P.Chartrand, submitted). We propose that Loc1p’s role in the nuclear export of *ASH1* mRNA is to increase the efficiency of *ASH1* mRNA recognition by the cytoplasmic

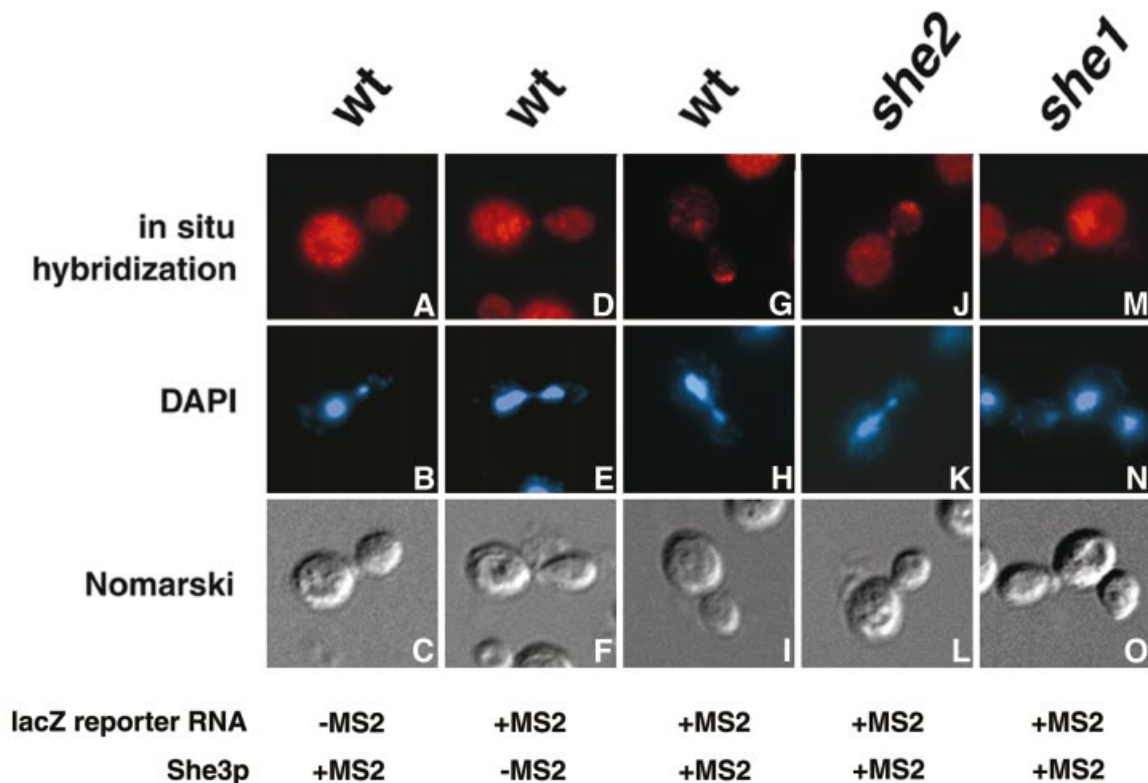


Fig. 6. RNA localization directed by She3p-MS2 is independent of She2p. (A–C) Representative images for *in situ* hybridization to lacZ mRNA, DAPI staining and Nomarski optics of wild-type cells expressing the She3p-MS2 fusion protein and the lacZ reporter RNA devoid of MS2 binding sites. (D–F) Images for wild-type cells expressing She3p and the lacZ-MS2 reporter RNA. (G–I) Images for wild-type cells expressing the She3p-MS2 fusion protein and the lacZ-MS2 reporter mRNA. Images for *she2* (J–L) and *she1* (M–O) cells, respectively, expressing the She3p-MS2 fusion protein and the lacZ-MS2 reporter mRNA.

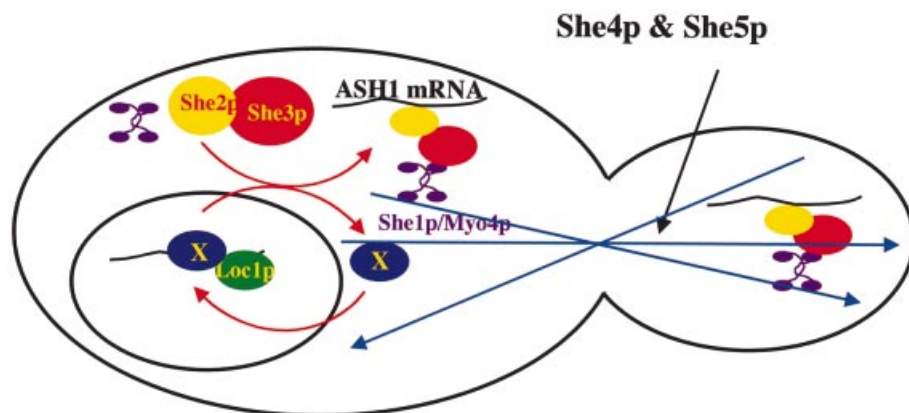


Fig. 7. Model for the localization of *ASH1* mRNA. In the nucleus, *ASH1* mRNA is identified by Loc1p in association with nucleocytoplasmic shuttling proteins. Following transport through the nuclear pores, *ASH1* mRNA is rapidly assembled into a transport particle through the RNA-binding activity of She2p. The She2p-*ASH1* mRNA complex associates with Myo4p-She3p to transport *ASH1* mRNA to the bud tip by movement on actin cables. Once at the bud tip, *ASH1* mRNA will be anchored through a mechanism that remains to be determined.

localization machinery. A component of the cytoplasmic localization machinery, perhaps She2p, might shuttle between the cytoplasm and nucleus to receive *ASH1* mRNA from Loc1p. Following transport through the nuclear pore, *ASH1* mRNA would be rapidly identified and bound by the cytoplasmic RNA localization machinery: She1p, She2p and She3p.

She2p's sole function appears to be to associate *ASH1* mRNA indirectly to the motor protein Myo4p, essential for

the proper localization of *ASH1* mRNA (Long *et al.*, 1997; Takizawa *et al.*, 1997) and an *ASH1* mRNA-GFP particle (Bertrand *et al.*, 1998; Beach *et al.*, 1999). Myo4p and She3p have been found to immunoprecipitate with *ASH1* mRNA. (Münchow *et al.*, 1999; Takizawa and Vale, 2000). This *ASH1* mRNA-Myo4p association requires both She2p and She3p. Moreover, Myo4p and She3p can interact in a two-hybrid interaction assay and co-immunoprecipitate (Münchow *et al.*, 1999; Takizawa

Table I. Yeast strains used in this study

Strain	Genotype	Source
YBZ-1	<i>Mata, ura3-52, leu2-3,112, his3-200, trp1-1, ade2, LYS2:::(LexAop)-lacZ, LexA-MS2 -MS2 coat (N55K)</i>	M.Wickens
PJ69-4a	<i>Mata, trp1-901, leu2-3,112, ura3-52, his3-200, gal4, gal80, GAL2-ADE2, LYS2:::GAL1-HIS3, met2:::GAL7-lacZ</i>	James <i>et al.</i> (1996)
K699	<i>Mata, ura3, leu2-3,112, his3-11, trp1-1, ade2-1, ho, can1-100</i>	Jansen <i>et al.</i> (1996)
YLM195	K699 <i>she1::KAN</i>	this study
YLM197	K699 <i>she2::KAN</i>	this study
YLM445	PJ69-4a <i>ash1::KAN</i>	this study
YLM584	YBZ-1 <i>she1::KAN</i>	this study
YLM585	YBZ-1 <i>she2::KAN</i>	this study
YLM586	YBZ-1 <i>she3::KAN</i>	this study
YLM587	YBZ-1 <i>she4::KAN</i>	this study
YLM588	YBZ-1 <i>she5::KAN</i>	this study

and Vale, 2000). We postulate that She2p functions to recruit the Myo4p–She3p complex to the four localization elements in the *ASH1* mRNA sequence. Altogether, these results point to a role of She3p in bridging the interaction of *ASH1* mRNA with the transport mechanism, specifically Myo4p. Our results support this hypothesis by providing direct evidence of RNA-dependent interactions between She2p and *ASH1* mRNA. Furthermore, She2p is unnecessary for She3p-dependent RNA localization when She3p is fused to the MS2 RNA-binding protein. This observation is consistent with the hypothesis that She2p's sole function in *ASH1* mRNA localization is to engage Myo4p–She3p with *ASH1* mRNA.

Finally, this RNA localization complex would transport and localize the *ASH1* mRNA to the distal tip of daughter cells using polarized actin filaments. Organization of the actin cytoskeleton requires She5p and probably She4p. It remains to be determined how *ASH1* mRNA is anchored at the bud tip upon reaching its destination.

Materials and methods

Growth media, yeast strains and plasmids

Yeast cells were grown in either synthetic growth media lacking the nutrients indicated or rich media (Rose *et al.*, 1990). Yeast strains used in this study are listed in Table I. Yeast gene disruption cassettes were created by PCR amplification of the *loxP-KAN-loxP* construct in plasmid pUG6 and primers specific for the gene of interest (Güldener *et al.*, 1996).

Yeast cells transformed with a *KAN* disruption cassette were plated on YEPD plates containing 200 µg/ml G418. The original selection plates were replica plated to fresh YEPD + G418 plates, and positive colonies were isolated. Genomic DNA was extracted from the candidate colonies using the DNA-Pure Yeast Genomic Kit (CPG Inc.). Specific disruptions were confirmed by PCR analysis of genomic DNA.

Plasmids used for these studies were constructed using standard techniques. The DNA fragments containing the sequences of elements E1 (153 bp), E2A (153 bp) and E3 (127 bp from 2741 to 2867) were cloned in the *SmaI* site of plasmid pIII/MS2-2, generating plasmids pXR196, pXR197 and pRL80, respectively. The specific mutants in element E3 (mutants M9, M9 + 9A, M13 and M14; Chartrand *et al.*, 1999) were also cloned into the *SmaI* site of pIII/MS2-2 (see Table II).

Plasmids pEP13 and pRL174 were generated by cloning the ORFs corresponding to She3p and She2p into plasmids pACT2 and pGBDU-c2, respectively.

Plasmid pRL172 was created by cloning the ORF corresponding to She2p into plasmid pGEX-2T, respectively.

Plasmids pRL168, pRL176, pRL177, pRL179 and pRL180 correspond to plasmid pGEM-3Z containing the indicated *ASH1* fragments cloned as *BamHI-PstI* fragments. Plasmid pRL168 corresponds to nt 1750–1867 (E3 localization element); pRL176 corresponds to nt 598–750 (E1 localization element); pRL177 corresponds to nt 1044–1197 (E2A localization element); pRL179 corresponds to nt 1198–1446 (E2B); and

pRL180 corresponds to nt 1–1867 (*ASH1* ORF plus E3 element). Plasmids pRL194, pRL195, pRL196 and pRL197 contain the M9, M9 + M9A, M13 and M14 mutation element E3, respectively.

Plasmids pRL128 and pRL141 express She3p and She3p–MS2, respectively. Plasmid pRL128 was constructed by subcloning PCR products of *SHE3* into YEplac112, resulting in a *BamHI* restriction site immediately preceding the *SHE3* termination codon. A PCR product corresponding to the MS2 RNA-binding protein was subcloned into plasmid pRL128 as a *BamHI* fragment, generating plasmid pRL141.

Three-hybrid screen

The components for the three-hybrid selection were generously provided by the laboratory of M.Wickens, and the selection was performed essentially as described elsewhere (SenGupta *et al.*, 1996; Zhang *et al.*, 1997). *HIS3* and *lacZ* serve as reporter genes for the three-hybrid assay. Briefly, yeast strain YBZ-1 containing plasmid pRL80 was transformed with an *S.cerevisiae* cDNA activation library prepared in the S.Elledge laboratory (Li *et al.*, 1994) and obtained from the American Type Culture Collection (ATCC). Transformants were selected on media lacking leucine and histidine and containing 6.67 mg/l adenine and 1 mM 3-aminotriazole (3-AT). The three-hybrid assay contains a screen for RNA dependence. The plasmid expressing the fusion RNA also contains *URA3* and *ADE2*. When the yeast strain YBZ-1 is transformed with this plasmid and grown under non-selective conditions, this plasmid can be lost from the yeast cells, resulting in red and red/white sector colonies. When grown on the selection media lacking leucine and histidine and containing 6.67 mg/l adenine and 1 mM 3-AT, colonies that remain completely white represent candidates that require the fusion RNA for *HIS3* expression. Approximately 5×10^5 total yeast transformants were screened. After 1 week at 30°C, 49 all white colonies were observed. From these 49 colonies 23 were observed to express β-galactosidase. By selection for the loss of the fusion RNA plasmid by plating on media containing 5-fluoroorotic acid, only 9 out of 23 expressed *HIS3* and β-galactosidase dependent on the hybrid RNA. The cDNA inserts for these clones were determined, and one of the clones corresponded to She3p. In a subsequent three-hybrid screen with 2.7×10^6 transformants, one of the clones meeting all criteria corresponded to She3p.

Two-hybrid analysis

Yeast strain PJ69-4a was transformed with the following combinations of plasmids: pRL174 pEP13, pRL174 pACT2, pGBDU-C2 pEP13 and pGBDU-C2 pACT2. Yeast transformants were assayed for *HIS3* reporter gene expression by assaying growth on synthetic 2% glucose, 1 mM 3-AT media lacking leucine, uracil and histidine. *LacZ* expression was determined using *O*-nitrophenyl-β-D-galactopyranoside (ONPG).

LacZ expression

Qualitative β-galactosidase expression levels were determined by X-gal filter assay and quantitative β-galactosidase expression levels were determined by liquid assay using ONPG (Guarente, 1983). The normalized β-galactosidase expression levels represent the average of three independent experiments.

In vitro RNA transcription

For UV cross-linking assays, plasmids pRL168, pRL176, pRL179, pRL194, pRL195, pRL196 and pRL197 were linearized by digestion with *HindIII*, while plasmid pRL177 was linearized with *PstI*. *In vitro*

Table II. Plasmids used in this study

Plasmid	Features	Source
pACT2	yeast vector for expressing Gal4p activation domain fusion proteins	Li <i>et al.</i> (1994)
pGBDU-c2	yeast vector for expression Gal4p DNA binding domain fusion proteins	James <i>et al.</i> (1996)
pGEX-2T	<i>Escherichia coli</i> vector for generating GST-tagged fusion proteins	Pharmacia
YEplac195	yeast multicopy shuttle plasmid marked with URA3	Gietz and Sugino (1998)
YEplac195-lacZ-ADHII	LacZ coding sequence with ADHII 3'-UTR cloned in plasmid YEplac195	Chartrand <i>et al.</i> (1999)
YEplac195-lacZ-MS2-ADHII	Yeplac195-lacZ-ADHII with six MS2 stem-loop motifs between lacZ and ADHII	Bertrand <i>et al.</i> (1998)
YEplac112	yeast multicopy shuttle plasmid marked with LEU2	Gietz and Sugino (1998)
pGEM-3Z	vector for <i>in vitro</i> transcription	Promega
pGEM-4Z	vector for <i>in vitro</i> transcription	Promega
pUG6	plasmid for generating KAN disruption cassettes by PCR	J.H.Hegemann
pIIIA/MS2-2	three-hybrid vector for the expression of MS2 fusion RNAs	M.Wickens
pIII/IRE-MS2	three-hybrid plasmid expressing MS2-IRE fusion RNA	M.Wickens
pEP13	pACT2 containing entire She3p ORF	this work
pRL80	pIIIA/MS2-2 containing the 127 nt <i>Sma</i> I E3 <i>ASH</i> I localization element	this work
pRL128	Yeplac112 containing <i>SHE</i> 3	this work
pRL141	Yeplac112 expressing She3p-MS2	this work
pRL168	pGEM-3Z containing element E3 (118 nt)	this work
pRL172	pGEX-2T containing the ORF corresponding to She2p	this work
pRL176	pGEM-3Z containing element E1 (153 nt)	this work
pRL177	pGEM-3Z containing element E2A (154 nt)	this work
pRL179	pGEM-3Z containing element E2B (249 nt)	this work
pRL180	pGEM-3Z containing elements E1, E2A, E2B and E3 (1868 nt)	this work
pRL174	pGBDU-c2 containing She2p ORF	this work
pRL194	element E3-mutant M9 (118 nt) cloned in pGEM-3Z	this work
pRL195	element E3-mutant M9 + 9A (118 nt) cloned in pGEM-3Z	this work
pRL196	element E3-mutant M13 (118 nt) cloned in pGEM-3Z	this work
pRL197	element E3-mutant M13 (118 nt) cloned in pGEM-3Z	this work
pRL204	pIIIA/MS-2 containing 130 bp from the ADHII 3'-UTR	this work
pRL205	pACT plasmid containing She3p clone (amino acids 236–425) isolated through three-hybrid analysis	this work
pXR64	Yeplac195-lacZ-ADHII containing element E3 (118 nt)	Chartrand <i>et al.</i> (1999)
pXR113	Yeplac195-lacZ-ADHII containing element E1 (150 nt)	Chartrand <i>et al.</i> (1999)
pXR137	Yeplac195-lacZ-ADHII containing element E2A (150 nt)	Chartrand <i>et al.</i> (1999)
pXR150	Yeplac195-lacZ-ADHII containing element E2B (275 nt)	Chartrand <i>et al.</i> (1999)
pXR192	element E3-mutant M9 (118 nt) cloned in pIIIA/MS2-2	this work
pXR193	element E3-mutant M9 + 9A (118 nt) cloned in pIIIA/MS2-2	this work
pXR194	element E3-mutant M13 (118 nt) cloned in pIIIA/MS2-2	this work
pXR195	element E3-mutant M14 (118 nt) cloned in pIIIA/MS2-2	this work
pXR196	element E1 (150 nt) cloned in pIIIA/MS2-2	this work
pXR197	element E2A (150 nt) cloned in pIIIA/MS2-2	this work

transcription reactions using linearized pRL168, pRL176, pRL177, pRL179, pRL180, pRL194, pRL195, pRL196 and pRL197 were performed using T7 RNA polymerase. ³²P-labeled RNAs were purified by PAGE on 6% denaturing gels. In the case of synthesis of unlabeled RNA, purification was achieved by DNase digestion, phenol/chloroform extraction and ethanol precipitation.

GST-She2p expression and purification

A single colony of *E.coli* DH5 α containing either pGEX-2T or pRL172 was inoculated into 50 ml of LB + amp, and the culture was grown overnight at 30°C. The overnight cultures were diluted 1:50 into fresh LB + amp medium prewarmed to 30°C, and cultures were grown to an OD₆₀₀ of 0.500. Cultures were induced with 1 mM isopropyl- β -D-thiogalactopyranoside and incubated at 30°C for an additional 2 h. Cells were harvested by centrifugation and resuspended in 1 \times phosphate-buffered saline containing 2 μ g/ml aprotinin, 2 μ g/ml leupeptin, 1 μ g/ml pepstatin and 175 μ g/ml phenylmethylsulfonyl fluoride. Extracts were prepared by passing the cell suspension three times through a French press, and the soluble fraction recovered after centrifugation at 31 000 g for 15 min at 4°C. The soluble fraction was aliquoted and frozen at -80°C. GST and GST-She2p were batch purified using GST-Sepharose according to the supplier's (Pharmacia) recommendations.

UV cross-linking assays

For UV cross-linking assays, recombinant protein and ³²P-labeled RNA were incubated for 10 min at 30°C in 20 mM HEPES pH 7.9, 33 mM KCl, 3.2 mM MgCl₂, 6% glycerol and 4 mM dithiothreitol in a 25 μ l reaction

volume containing the indicated amounts of protein. For competition studies, unlabeled RNAs (amounts are indicated in the figure legends) were added just prior to the addition of labeled RNAs to the recombinant proteins. Subsequently, reaction mixtures were UV cross-linked for 20 min at 4°C using a 254 nm UV lamp from Spectroline, model number EF-180C, at a distance of 9 cm. Subsequently, RNase A was added to 0.75 mg/ml, and reactions incubated at 37°C for 15 min. Following RNase treatment, 5 μ l of 6 \times sample buffer were added and the reactions were analyzed by SDS-PAGE. Gels were stained briefly with Coomassie Blue, dried and autoradiography performed.

In situ hybridization

Yeast cells were grown in the appropriate medium to early or mid-log phase, fixed with formaldehyde and spheroplasted as described previously (Long *et al.*, 1995). For *in situ* hybridization, yeast spheroplasts were hybridized with a pool of Cy3-conjugated lacZ DNA oligonucleotide probes following the protocol described elsewhere (Long *et al.*, 1995).

Acknowledgements

We would like to thank Ralf-Peter Jansen for communicating results prior to publication. We would also like to thank Marvin Wickens for assistance as well as providing the yeast strains and plasmids for the three-hybrid assay. This work was supported by NIH grant GM57071 to R.H.S., Canadian FCAR and CIHR Fellowships to P.C. and Training

Grant (2-T32-CA09475) to W.G. R.M.L. was supported by NIH grants F32 HD08088 and GM60392, the Pew Scholars Program in the Biomedical Sciences, the March of Dimes and the Medical College of Wisconsin Research Affairs Committee.

References

- Beach,D.L., Salmon,E.D. and Bloom,K. (1999) Localization and anchoring of mRNA in budding yeast. *Curr. Biol.*, **9**, 569–578.
- Bertrand,E., Chartrand,P., Schaefer,M., Shenoy,S.M., Singer,R.H. and Long,R.M. (1998) Localization of *ASH1* mRNA particles in living yeast. *Mol. Cell*, **2**, 437–445.
- Bobola,N., Jansen,R.-P., Shin,T.H. and Nasmyth,K. (1996) Asymmetric accumulation of Ash1p in postanaphase nuclei depends on a myosin and restricts yeast mating-type switching to mother cells. *Cell*, **84**, 699–709.
- Chartrand,P., Meng X., Singer,R.H. and Long,R.M. (1999) Structural elements required for the localization of *ASH1* mRNA and of a green fluorescent protein reporter particle *in vivo*. *Curr. Biol.*, **9**, 333–336.
- Evangelista,M., Blundell,K., Longtine,M.S., Chow,C.J., Adames,N., Pringle,J.R., Peter,M. and Boone,C. (1997) Bni1p, a yeast formin linking Cdc42p and the actin cytoskeleton during polarized morphogenesis. *Science*, **276**, 118–122.
- Fields,S. and Song,O. (1989) A novel genetic system to detect protein–protein interactions. *Nature*, **340**, 245–246.
- Fouts,D., True,H. and Celander,D. (1997) Functional recognition of fragmented operator site by R17/MS2 coat protein, a translational repressor. *Nucleic Acids Res.*, **25**, 4464–4473.
- Gietz,R.D. and Sugino,A. (1988) New yeast–*Escherichia coli* shuttle vectors constructed with *in vitro* mutagenized yeast genes lacking six-base pair restriction sites. *Gene*, **74**, 527–534.
- Gonzalez,I., Buonomo,S.B., Nasmyth,K. and von Ahsen,U. (1999) *ASH1* mRNA localization in yeast involves multiple secondary structural elements and Ash1 protein translation. *Curr. Biol.*, **9**, 337–340.
- Guarente,L. (1983) Yeast promoters and lacZ fusions designed to study expression of cloned genes in yeast. *Methods Enzymol.*, **101**, 181–191.
- Güldener,U., Heck,S., Fiedler,T., Beinhauer,J. and Hegemann,J.H. (1996) A new efficient gene disruption cassette for repeated use in budding yeast. *Nucleic Acids Res.*, **24**, 2519–2524.
- Haarer,R.K., Petzold,A., Lillie,S.H. and Brown,S.S. (1994) Identification of *MYO4*, a second class V myosin gene in yeast. *J. Cell Sci.*, **107**, 1055–1064.
- James,P., Halladay,J. and Craig,E.A. (1996) Genomic libraries and a host strain designed for highly efficient two-hybrid selection in yeast. *Genetics*, **144**, 1425–1436.
- Jansen,R.-P., Dowser,C., Michaelis,C., Galova,M. and Nasmyth,K. (1996) Mother cell-specific *HO* expression in budding yeast depends on the unconventional myosin Myo4p and other cytoplasmic proteins. *Cell*, **84**, 687–697.
- Klausner,R.D., Rouault,T.A. and Harford,J.B. (1993) Regulating the fate of mRNA: the control of cellular iron metabolism. *Cell*, **72**, 19–28.
- Kohno,H. *et al.* (1996) Bni1p implicated in cytoskeletal control is a putative target of Rho1p small GTP binding protein in *Saccharomyces cerevisiae*. *EMBO J.*, **15**, 6060–6068.
- Li,L., Elledge,S.J., Peterson,C.A., Bales,E.S. and Legerski,R.J. (1994) Specific association between the human DNA repair proteins XPA and ERCC1. *Proc. Natl Acad. Sci. USA*, **91**, 5012–5016.
- Long,R.M., Elliott,D.J., Stutz,F., Rosbash,M. and Singer,R.H. (1995) Spatial consequences of defective processing of specific yeast mRNAs revealed by fluorescent *in situ* hybridization. *RNA*, **1**, 1071–1078.
- Long,R.M., Singer,R.H., Meng,X., Gonzalez,I., Nasmyth,K. and Jansen,R.-P. (1997) Mating type switching in yeast controlled by asymmetric localization of *ASH1* mRNA. *Science*, **277**, 383–387.
- Meleforts,O. and Hentze,M.W. (1993) Translational regulation by mRNA/protein interactions in eukaryotic cells: ferritin and beyond. *BioEssays*, **15**, 85–90.
- Münchow,S., Sauter,C. and Jansen,R.-P. (1999) Association of the class V myosin Myo4p with a localised messenger RNA in budding yeast depends on She proteins. *J. Cell Sci.*, **112**, 1511–1518.
- Nasmyth,K. and Jansen,R.-P. (1997) The cytoskeleton in mRNA localization and cell differentiation. *Curr. Opin. Cell Biol.*, **3**, 396–400.
- Oleynikov,Y. and Singer,R.H. (1998) RNA localization: different zipcodes, same postman? *Trends Cell Biol.*, **8**, 381–383.
- Rose,M.D., Winston,F. and Hieter,P. (1990) *Methods in Yeast Genetics. A Laboratory Course Manual*. Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY.
- Schnorrer,F., Bohmann,K. and Nusslein-Volhard,C. (2000) The molecular motor dynein is involved in targeting swallow and bicoid mRNA to the anterior pole of *Drosophila* oocytes. *Nature Cell Biol.*, **2**, 185–190.
- SenGupta,D.J., Zhang,B., Kraemer,B., Pochart,P., Field,S. and Wickens,M. (1996) A three-hybrid system to detect RNA–protein interactions *in vivo*. *Proc. Natl Acad. Sci. USA*, **93**, 8496–8501.
- Sil,A. and Herskowitz,I. (1996) Identification of asymmetrically localized determinant, Ash1p, required for lineage-specific transcription of the yeast *HO* gene. *Cell*, **84**, 711–722.
- Takizawa,P. and Vale,R. (2000) The myosin motor, Myo4p, binds Ash1 mRNA via the adapter protein, She3p. *Proc. Natl Acad. Sci. USA*, **97**, 5273–5278.
- Takizawa,P., Sil,A., Swedlow,J., Herskowitz,I. and Vale,R. (1997) Actin-dependent localization of an RNA encoding a cell-fate determinant in yeast. *Nature*, **389**, 90–93.
- Wendland,B., McCaffery,J.M.N., Qin,X. and Emr,S.D. (1996) A novel fluorescence-activated cell sorter-based screen for yeast endocytosis mutants identifies a yeast homologue of mammalian eps15. *J. Cell Biol.*, **135**, 1485–1500.
- Zhang,B., Gallegos,M., Puoti,A., Durkin,E., Field,S., Kimble,J. and Wickens,M.P. (1997) A conserved RNA-binding protein that regulates sexual fates in the *C.elegans* hermaphrodite germ line. *Nature*, **390**, 477–484.

Received July 5, 2000; revised October 10, 2000;
accepted October 13, 2000