

# CKI $\alpha$ Is Associated with and Phosphorylates Star-PAP and Is Also Required for Expression of Select Star-PAP Target Messenger RNAs\*

Received for publication, January 25, 2008, and in revised form, February 20, 2008. Published, JBC Papers in Press, February 27, 2008, DOI 10.1074/jbc.M800656200

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We have recently identified Star-PAP, a nuclear poly(A) polymerase that associates with phosphatidylinositol-4-phosphate 5-kinase I $\alpha$  (PIPKI $\alpha$ ) and is required for the expression of a specific subset of mRNAs. Star-PAP activity is directly modulated by the PIPKI $\alpha$  product phosphatidylinositol 4,5-bisphosphate (PI-4,5-P<sub>2</sub>), linking nuclear phosphoinositide signaling to gene expression. Here, we show that PI-4,5-P<sub>2</sub>-dependent protein kinase activity is also a part of the Star-PAP protein complex. We identify the PI-4,5-P<sub>2</sub>-sensitive casein kinase I $\alpha$  (CKI $\alpha$ ) as a protein kinase responsible for this activity and further show that CKI $\alpha$  is capable of directly phosphorylating Star-PAP. Both CKI $\alpha$  and PIPKI $\alpha$  are required for the synthesis of some but not all Star-PAP target mRNA, and like Star-PAP, CKI $\alpha$  is associated with these messages *in vivo*. Taken together, these data indicate that CKI $\alpha$ , PIPKI $\alpha$ , and Star-PAP function together to modulate the production of specific Star-PAP messages. The Star-PAP complex therefore represents a location where multiple signaling pathways converge to regulate the expression of specific mRNAs.

Polyadenylation of most mRNAs is required for their efficient transcription and export as well as regulating their stability and translational efficiency (1). Polyadenylation of mRNA is achieved through the activity of poly(A) polymerases (PAPs).<sup>2</sup> There are multiple PAPs in mammalian cells, including PAP $\alpha$ , which is thought to be primarily responsible for the polyadenylation of newly transcribed mRNAs in the nucleus (2, 3). Additionally, there are “non-canonical” PAPs, including Gld2 and Trf4, which regulate stability and degradation of specific RNAs through polyadenylation (4, 5).

Star-PAP is a non-canonical poly(A) polymerase that is distinct from all other currently characterized PAP enzymes (6). Like the canonical PAP $\alpha$ , Star-PAP has an RNA recognition motif and is a nuclear enzyme. However, similar to non-canon-

ical PAPs, the Star-PAP protein complex and architecture differ significantly from PAP $\alpha$ , and consequently, Star-PAP specifically targets a select subset of mRNAs. This suggests that Star-PAP is a hybrid PAP that is required for the 3'-end formation of newly transcribed pre-mRNAs but functions in a regulatory role to control mRNA expression levels. Star-PAP has a unique domain structure relative to all other known PAPs (6). One unique feature is a 205-amino acid proline-rich region (PRR) inserted into the catalytic PAP core. The PRR splits the catalytic PAP domain, and this region represents a potential site for regulation of Star-PAP function.

Poly(A) polymerases operate as large multiprotein complexes responsible for the 3'-processing of RNAs (7, 8). The Star-PAP polyadenylation complex is similar to that of poly(A) polymerases that utilizes mRNA as a substrate and includes cleavage and polyadenylation specificity factor (CPSF) subunits, cleavage stimulatory factor (CstF) subunits, symplekin, and RNA polymerase II (6, 9). However, unlike other poly(A) polymerase complexes, the Star-PAP complex also contains phosphatidylinositol-4-phosphate 5-kinase I $\alpha$  (PIPKI $\alpha$ ). Moreover, Star-PAP poly(A) polymerase activity is stimulated by the PIPKI $\alpha$  product phosphatidylinositol 4,5-bisphosphate (PI-4,5-P<sub>2</sub>). Functionally, PIPKI $\alpha$  is required for the expression of select Star-PAP target mRNAs (6).

Phosphoinositides, including PI-4,5-P<sub>2</sub>, are lipid signaling molecules that act as important regulators of numerous cellular functions (10). Phosphoinositides are generated ubiquitously in cells by multiple families of phosphoinositide kinases, which include PIPKI $\alpha$  (11). Phosphoinositide signaling specificity is based on the spatial restriction of these kinases to specific subcellular locations (12). This is often achieved through specific interactions between the phosphoinositide-generating enzymes and targeting factors, which are themselves often PI-4,5-P<sub>2</sub>-regulated proteins (13, 14). This allows for the generation of inositol lipid second messengers at specific sites within the cell that, in turn, allows for precise targeting of individual phosphoinositide-sensitive pathways (12).

In the nucleus, PI-4,5-P<sub>2</sub> and PIPKI $\alpha$ , as well as Star-PAP, are found in structures called nuclear speckles (6, 15), which are nuclear bodies enriched in factors required for the processing of pre-mRNA (16). The direct interaction between Star-PAP and PIPKI $\alpha$  and the proximity of Star-PAP to PI-4,5-P<sub>2</sub> generation in nuclear speckles suggest that nuclear phosphoinositides are spatially positioned to modulate Star-PAP activity *in vivo*. The presence of PIPKI $\alpha$  in the Star-PAP complex implies

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<sup>2</sup> The abbreviations used are: PAP, poly(A) polymerase; PRR, proline-rich region; CPSF, cleavage and polyadenylation specificity factor; CstF, cleavage stimulatory factor; siRNA, small interfering RNA; MBP, myelin basic protein; tBHQ, *tert*-butylhydroquinone; CKI, casein kinase I; PI-4,5-P<sub>2</sub>, phosphatidylinositol 4,5-bisphosphate; PIPKI $\alpha$ , phosphatidylinositol-4-phosphate 5-kinase I $\alpha$ .

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that phosphoinositide-based signal transduction pathways may be able to modulate Star-PAP function.

Another PI-4,5-P<sub>2</sub>-sensitive protein found at nuclear speckles is the protein kinase CKI $\alpha$  (formerly known as casein kinase I $\alpha$ ) (17). The protein kinase activity of CKI $\alpha$  is specifically inhibited by PI-4,5-P<sub>2</sub> (18). CKI $\alpha$  is a member of the ubiquitously expressed CKI family of constitutively active Ser/Thr-specific protein kinases (19). CKI activity is modulated through the unique recognition motif (S/T)(P)XX(S/T), where S/T represents a phosphoserine or phosphothreonine in the -3 position (20). The activity of CKIs toward their substrates is regulated through this "priming" phosphorylation that precedes phosphorylation by CKIs. However, this is not the only mechanism of regulation for CKI family members.

Because the composition of a polyadenylation complex is important for the function of the associated poly(A) polymerase (5, 9), we looked for additional unique components of the Star-PAP-associated complex relative to PAP $\alpha$ . Identification of other unique Star-PAP-associated factors will lead to a more complete understanding of Star-PAP function and of its differences from other known PAPs. Here, we show that the Star-PAP complex contains unique protein kinase activities compared with PAP $\alpha$ . One of these kinases is the PI-4,5-P<sub>2</sub>-sensitive kinase CKI $\alpha$ , which is capable of directly phosphorylating Star-PAP. Moreover, CKI $\alpha$  and PIPKI $\alpha$  are required for the expression of a subset of Star-PAP target mRNAs. These data further reinforce the existence of a nuclear phosphoinositide-based signaling complex that regulates the Star-PAP polyadenylation complex to modulate expression of specific mRNAs.

### EXPERIMENTAL PROCEDURES

**Cell Culture, Transfection, and Treatments**—HEK 293 and normal rat kidney cells were obtained from American Type Culture Collection and maintained in Dulbecco's modified Eagle's medium plus 10% fetal bovine serum at 37 °C in 5% CO<sub>2</sub>. For DNA transfection, cells were transfected using the calcium phosphate method with the indicated amounts of DNA. The growth medium was exchanged after 4 h, and the cells were harvested at the indicated time. siRNA oligonucleotides were transfected using calcium phosphate at a final concentration of 120 nM oligonucleotide/ml of growth medium. Growth medium was replaced 6 h after transfection, and the transfection was repeated 24 h later. Cells were harvested for analysis 72 h after the first transfection. For PIPKI $\alpha$ , the oligonucleotides used were PIPKI $\alpha$ -1 (GGUGCCAUCCAGUUAGGCA) and PIPKI $\alpha$ -3 (GAAGUUGGAGCACUCUUGG). For CKI $\alpha$ , an siGENOME SMARTpool (Dharmacon) was directed against CSNK1A1. To induce a transcriptional antioxidant response, HEK 293 cells were treated with 100  $\mu$ M *tert*-butylhydroquinone (tBHQ; Sigma) in Me<sub>2</sub>SO for 4 h. Control cells were treated with Me<sub>2</sub>SO only. CKI inhibitors CKI-7 (Sigma) and IC261 (Calbiochem) were resuspended in Me<sub>2</sub>SO and used at the final concentrations indicated.

**Immunofluorescence**—Normal rat kidney cells were plated on glass coverslips in 35-mm dishes. Cells were transfected with 2.5  $\mu$ g of FLAG-Star-PAP and allowed to express for 24 h. Coverslips were washed in phosphate-buffered saline and fixed for

10 min in -70 °C methanol. Immunofluorescence staining and microscopy were performed as described previously (13) using anti-FLAG M2 (Sigma-Aldrich) and rabbit anti-CKI $\alpha$  polyclonal antibodies.

**Expression and Affinity Purification of FLAG Proteins**—Human Star-PAP and rat CKI $\alpha$  cDNAs were cloned into the pFLAG-1 mammalian expression vector (Sigma). The kinase-dead CKI $\alpha$  mutant K46R was generated by PCR-based mutagenesis using primers 5'-gaagtggcagtgagactagaatcccag-3' and 5'-ctgggattctagtctcactgcccactcc-3'. For each FLAG purification,  $\sim 5 \times 10^6$  HEK 293 cells in four 10-cm dishes were transfected with 10  $\mu$ g of DNA and allowed to express for 48 h. FLAG purifications were performed as described previously (6) according to the manufacturer's instructions.

**Generation and Purification of Star-PAP Truncation Mutants**—FLAG-tagged Star-PAP phosphorylation site point mutants were generated using PCR-based mutagenesis. The wild-type and mutant constructs were transfected into HEK 293 cells, allowed to express for 24 h, and lysed in FLAG lysis buffer (6). 2  $\mu$ g of anti-FLAG M2 antibody was added to the clarified lysates and incubated at 4 °C with rotation for 1 h, followed by additional 1 h with protein A-Sepharose beads (Amersham Biosciences). The beads were washed three times with FLAG lysis buffer and resuspended in 1 $\times$  kinase buffer (50 mM Tris-HCl (pH 7.5), 10 mM MgCl<sub>2</sub>, and 0.5 mM EGTA) for use in kinase assays.

**Protein Kinase Assays**—Protein kinase assays were performed in 1 $\times$  kinase buffer. Assays were initiated by the addition of 10  $\mu$ M ATP and 5  $\mu$ Ci of [ $\gamma$ -<sup>32</sup>P]ATP to the reaction mixture. The endogenous kinase activity in the Star-PAP complex was destroyed by heating for 15 min at 65 °C. For inhibitor studies, all reaction components except ATP were incubated with inhibitors for 45 min on ice prior to starting the assay. CKI inhibitors IC261 (IC<sub>50</sub> = 11  $\mu$ M) and CKI-7 (IC<sub>50</sub>  $\sim$  6.0  $\mu$ M) were resuspended in Me<sub>2</sub>SO and used at final concentrations of 0.1–100  $\mu$ M. Synthetic PI-4,5-P<sub>2</sub> (Echelon Biosciences Inc.) was resuspended in 50 mM Tris-HCl (pH 7.9) at 2.5 mM, subjected to bath sonication to form micelles, and used at final concentrations of 12.5–100  $\mu$ M.

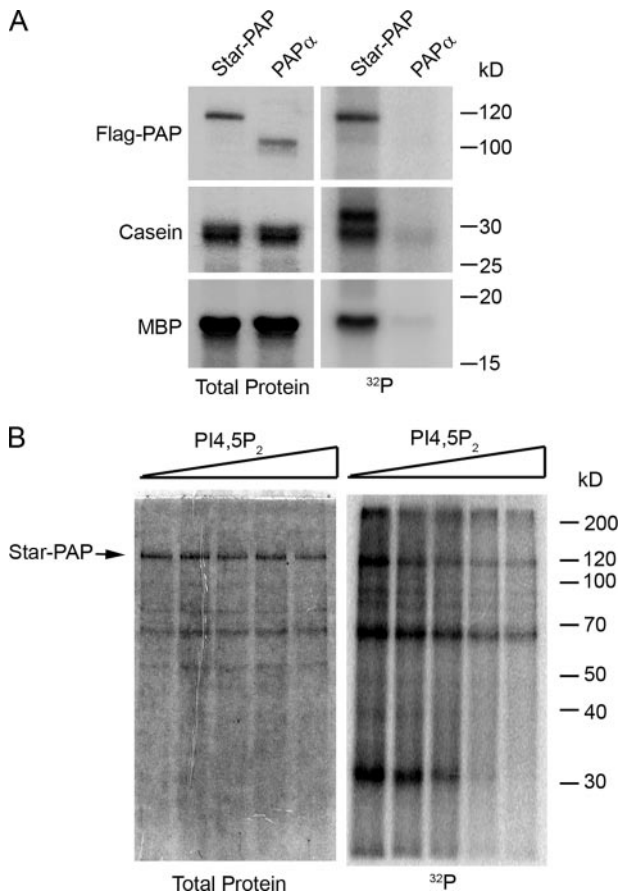
**Immunoprecipitation**—This was performed as described previously (6).

**Quantitative Real-time PCR**—This was performed as described previously (6). All mRNA levels were normalized to glyceraldehyde-3-phosphate dehydrogenase mRNA levels.

**RNA Immunoprecipitation**—This was performed as described previously (6).

### RESULTS

**The Star-PAP Complex Contains Protein Kinase Activity**—Purification of FLAG-Star-PAP or FLAG-PAP $\alpha$  from HEK 293 cells resulted in the copurification of a large protein complex (6). This complex contained factors essential for the 3'-end formation of mRNAs, including CPSF<sup>73</sup>, CPSF<sup>100</sup>, CstF<sup>64</sup>, symplekin, and RNA polymerase II (6). In contrast to PAP $\alpha$ , the Star-PAP complex also contained PIPKI $\alpha$  and, accordingly, PI-4-P kinase activity. To enhance the characterization of the Star-PAP complex, we subjected both the purified Star-PAP and



**FIGURE 1. Star-PAP is associated with PI-4,5-P $_2$ -sensitive kinase activity.** FLAG-Star-PAP or PAP $\alpha$  was expressed in HEK 293 cells, purified by anti-FLAG M2 affinity chromatography, and eluted in three consecutive fractions with a 3 $\times$ FLAG peptide. *A*, fractions were collected and used in an *in vitro* kinase assay with no substrate (*upper panels*), 100  $\mu$ g/ml casein (*middle panels*), or MBP (*lower panels*). The arrow indicates Star-PAP protein. *B*, the FLAG-Star-PAP complex was incubated with 0, 12.5, 25, 50, or 100  $\mu$ M PI-4,5-P $_2$  micelles for 45 min on ice prior to initiation of the kinase reaction by the addition of ATP.

PAP $\alpha$  complexes to an *in vitro* protein kinase assay to detect the presence of any associated protein kinases.

FLAG-Star-PAP and FLAG-PAP $\alpha$  were expressed in HEK 293 cells and purified on anti-FLAG M2 resin. Purified PAP complexes were subjected to an *in vitro* protein kinase assay with the generic protein kinase substrate myelin basic protein (MBP) or casein (100  $\mu$ g/ml). The purified Star-PAP complex contained protein kinase activity toward both MBP and casein, whereas the PAP $\alpha$  complex contained almost no detectable protein kinase activity (Fig. 1A). Interestingly, in these same assays, FLAG-Star-PAP was robustly phosphorylated (Fig. 1A), indicating that the associated kinase(s) can phosphorylate Star-PAP itself. These results identify protein kinase activity as another unique feature of the Star-PAP protein complex compared with PAP $\alpha$ . Furthermore, the ability of this kinase activity to phosphorylate Star-PAP suggests that phosphorylation may play a role in Star-PAP function.

The Star-PAP complex includes PIPKI $\alpha$ , and the activity of Star-PAP itself is directly regulated by the PIPKI $\alpha$  product PI-4,5-P $_2$ . Because of the relationship between Star-PAP and PI-4,5-P $_2$ , it is possible that other components of the Star-PAP complex may be regulated by PI-4,5-P $_2$  as well. Therefore, the

sensitivity of Star-PAP phosphorylation by the associated kinase(s) to PI-4,5-P $_2$  was examined. Phosphorylation of FLAG-Star-PAP by the associated kinase(s) was inhibited by PI-4,5-P $_2$  at concentrations as low as 12.5  $\mu$ M (Fig. 1B), indicating that the associated kinase activity is indeed sensitive to PI-4,5-P $_2$ . Remarkably, this is in the same range of PI-4,5-P $_2$  concentrations that stimulate Star-PAP poly(A) polymerase activity (6), indicating that both protein kinase and Star-PAP activities are regulated by similar concentrations of PI-4,5-P $_2$ .

*The PI-4,5-P $_2$ -sensitive Kinase CKI $\alpha$  Is Associated with the Star-PAP Complex*—There are very few protein kinases known to be inhibited by PI-4,5-P $_2$ ; one of these is CKI $\alpha$  (17). Accordingly, immunoblotting of purified FLAG complexes with a CKI $\alpha$ -specific antibody showed that CKI $\alpha$  copurified specifically with Star-PAP and not with PAP $\alpha$  (Fig. 2A). In addition, immunoprecipitation of endogenous Star-PAP from HEK 293 cells resulted in coprecipitation of endogenous CKI $\alpha$  (Fig. 2B), demonstrating that these two proteins reside in the same complex *in vivo*. Both Star-PAP (6) and CKI $\alpha$  (17) have been reported to localize at nuclear speckles. Consequently, there was a strong colocalization between FLAG-Star-PAP and endogenous CKI $\alpha$  in the nuclei of normal rat kidney cells at nuclear speckles (Fig. 2C), demonstrating that *in vivo* Star-PAP and CKI $\alpha$  are present at the same sites within the nucleus and are therefore spatially positioned to interact with each other.

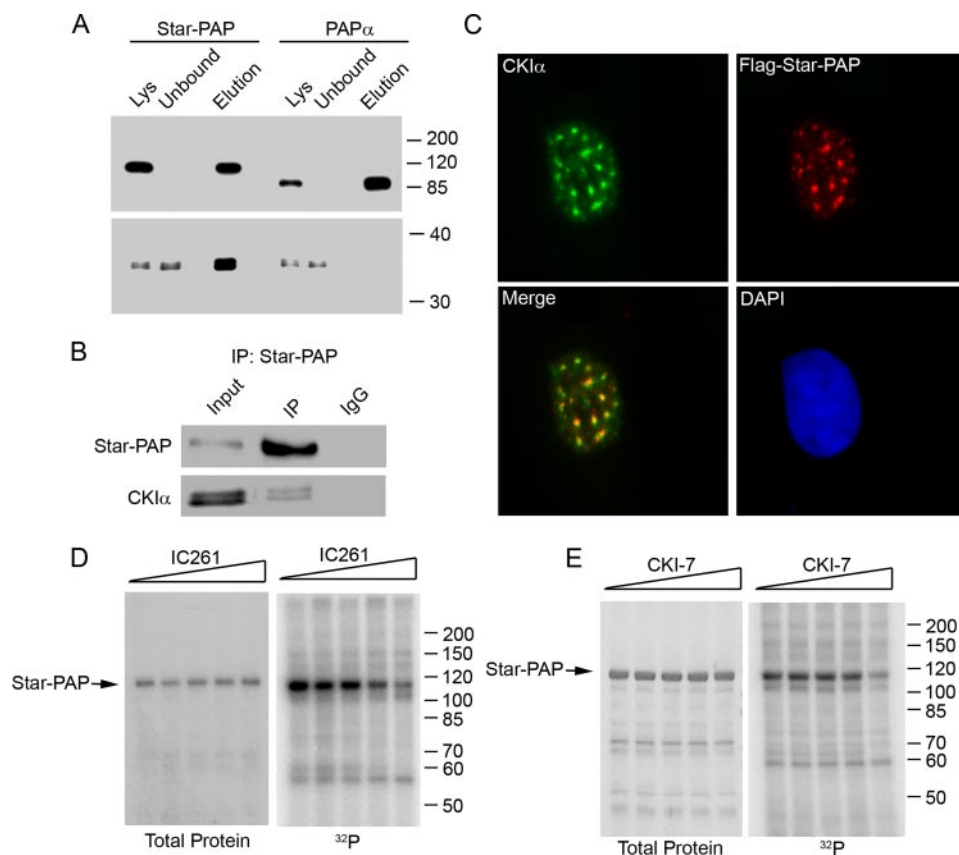
To confirm that the associated CKI $\alpha$  is involved in the phosphorylation of Star-PAP, the ability of CKI-specific inhibitors to block the phosphorylation of FLAG-purified Star-PAP by the associated kinase activity was examined. The CKI inhibitors IC261 (Fig. 2D) and CKI-7 (Fig. 2E) were both able to effectively block the phosphorylation of Star-PAP by the associated kinase activity in a dose-dependent fashion, signifying that CKI $\alpha$  is responsible for at least some of the kinase activity contained in the Star-PAP complex. The presence of CKI $\alpha$  specifically in the Star-PAP complex serves to further demonstrate that although Star-PAP and PAP $\alpha$  both contain the same basic 3'-end formation components, the Star-PAP complex contains additional signal transduction components that serve to distinguish it from the PAP $\alpha$  complex.

*CKI $\alpha$  Can Directly Phosphorylate Star-PAP in Vitro on the Proline-rich Insert Region*—The ability of IC261 and CKI-7 to block Star-PAP phosphorylation in the context of the purified complex suggests a role for CKI $\alpha$  in Star-PAP phosphorylation, but it does not prove that CKI $\alpha$  itself is directly phosphorylating Star-PAP. To demonstrate direct phosphorylation, purified CKI $\alpha$  was used to phosphorylate FLAG-purified Star-PAP. Prior to the assay, endogenous kinase activity in the FLAG complex was destroyed by heat inactivation. After heat inactivation, there was no detectable phosphorylation of Star-PAP (Fig. 3A). Purified FLAG-CKI $\alpha$  was able to directly phosphorylate heat-inactivated Star-PAP, whereas the catalytically inactive CKI $\alpha$  mutant K46R (21) was not (Fig. 3A). Similarly, phosphorylation by CKI $\alpha$  was blocked by 50  $\mu$ M IC261 or 50  $\mu$ M PI-4,5-P $_2$  (Fig. 3B). Together, these data indicate that the phosphorylation of Star-PAP is occurring directly by CKI $\alpha$ .

To determine the CKI $\alpha$  phosphorylation site(s) on Star-PAP, a series of FLAG-Star-PAP truncation and deletion mutants (Fig. 3C) was expressed and purified from HEK 293 cells and



## Phosphorylation of Star-PAP by CKI $\alpha$



**FIGURE 2. CKI $\alpha$  is associated with Star-PAP and is responsible for Star-PAP phosphorylation.** *A*, FLAG-purified Star-PAP and PAP $\alpha$  complexes were separated by SDS-PAGE, transferred to nitrocellulose, and probed with anti-FLAG and anti-CKI $\alpha$  antibodies. *B*, endogenous Star-PAP was immunoprecipitated (IP) from HEK 293 cells. The resulting precipitates were immunoblotted with Star-PAP- and CKI $\alpha$ -specific antibodies. A nonspecific IgG immunoprecipitation was used as a control. *C*, normal rat kidney cells were transfected with FLAG-Star-PAP, allowed to express for 24 h, and fixed for immunofluorescence. Cells were stained with anti-FLAG (red) and anti-CKI $\alpha$  (green) to determine subcellular localization. Colocalization is shown as yellow in the Merge panel. Nuclei are indicated by staining with 4',6-diamidino-2-phenylindole (DAPI). The purified FLAG-Star-PAP complex was incubated with 0, 0.1, 1.0, 10, or 100  $\mu$ M IC261 (IC<sub>50</sub> = 11  $\mu$ M) (*D*) or CKI-7 (IC<sub>50</sub> ~ 6.0  $\mu$ M) (*E*) prior to initiation of the kinase reaction by ATP. The arrow indicates Star-PAP protein.

subjected to the same *in vitro* kinase assays described above. CKI $\alpha$  was able to phosphorylate all truncation mutants except those that lacked the first half of the proline-rich region ( $\Delta$ PRR 1/2, amino acids 223–274) that splits the catalytic domain of Star-PAP (Fig. 3*D*), demonstrating that this region contains the CKI $\alpha$  phosphorylation site(s) in Star-PAP. This region contains nine serine and threonine residues conserved among mammalian species, including two consensus CKI $\alpha$  sites and a number of acidic residues that could contribute to additional CKI $\alpha$  phosphorylation sites (Fig. 3*E*).

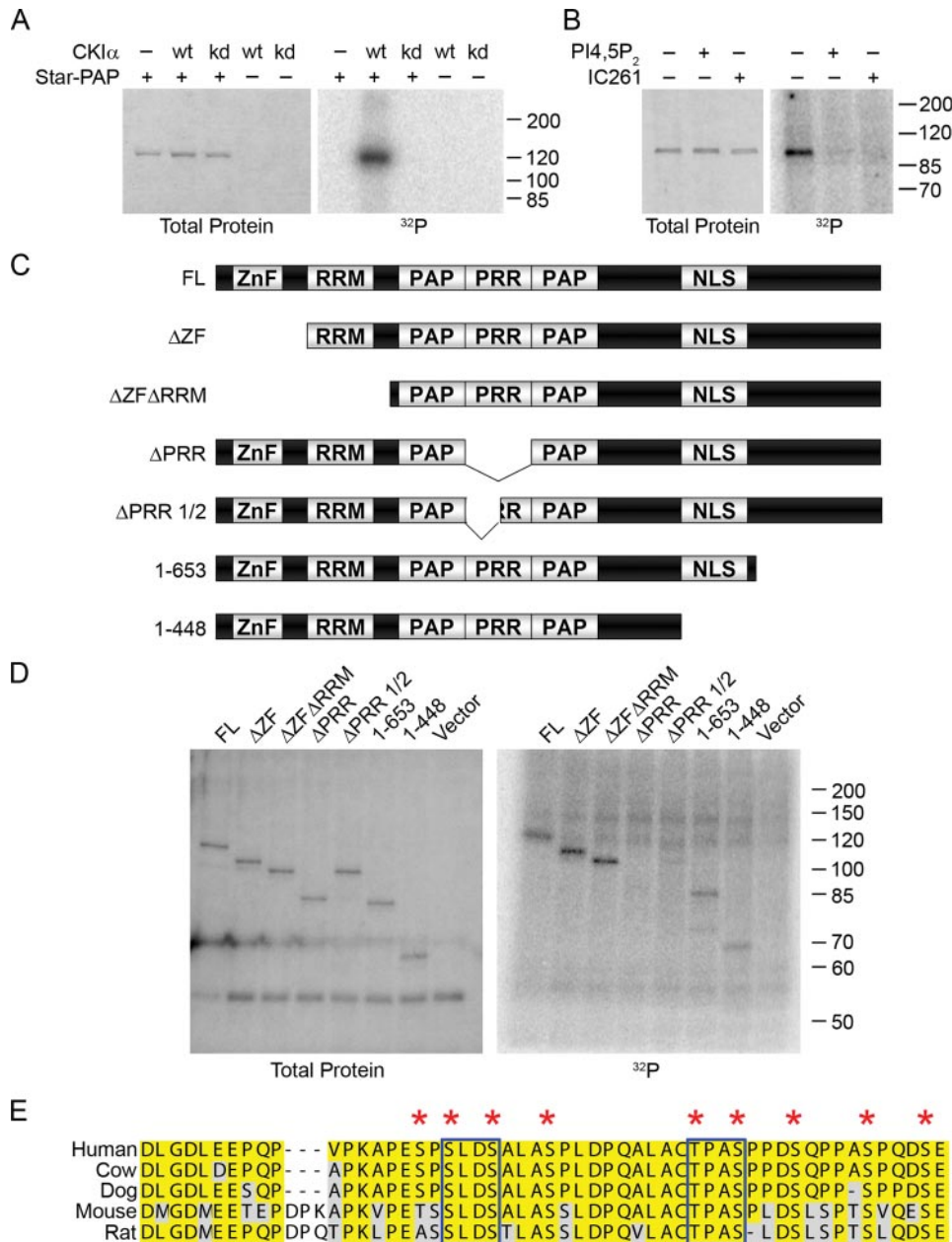
**The Star-PAP PRR Is Not Required for CKI $\alpha$  Association with the Star-PAP Complex**—The above results leave open the possibility that CKI $\alpha$  does not phosphorylate Star-PAP in the PRR but, rather, that CKI $\alpha$  requires the PRR to associate with the Star-PAP complex and subsequently phosphorylates other sites within Star-PAP. To examine this, the requirement for the Star-PAP PRR for association of CKI $\alpha$  and for protein kinase activity with the Star-PAP complex was assessed. Both full-length Star-PAP and Star-PAP $\Delta$ PRR expressed and purified from HEK 293 cells associated with endogenous CKI $\alpha$  (Fig. 4*A*). Furthermore, although FLAG-purified Star-PAP $\Delta$ PRR could not be phosphorylated by the associated kinases (Fig. 4*B*), the complex still

contained activity toward both casein and MBP similar to that of full-length Star-PAP (Fig. 4*C*), demonstrating that deletion of the PRR does not disrupt the association of protein kinase activity with the Star-PAP complex. This indicates that the inability of CKI $\alpha$  to phosphorylate Star-PAP $\Delta$ PRR mutants is most likely due to a deletion of the phosphorylation site(s) and not to a disruption of the Star-PAP/CKI $\alpha$  interaction.

**CKI $\alpha$  and PIPKI $\alpha$  Are Required for the Expression of Specific Star-PAP Target mRNAs**—Star-PAP and PIPKI $\alpha$  are required for expression of detoxifying and oxidative stress response mRNAs; however, microarray analysis indicated that Star-PAP is required for the expression of a diverse assortment of mRNAs (6). Therefore, the requirements for CKI $\alpha$  and PIPKI $\alpha$  for the expression of a diverse group of Star-PAP target mRNAs were analyzed. A group of mRNAs in which protein products function in a wide array of cellular processes was selected for further analysis, including heme oxygenase 1 (HO-1), NADPH:quinone oxidoreductase 1 (NQO1), cation transporter regulator-like 1 (CHAC1), asparagine synthetase (ASNS), p8 protein homolog (COM1), and secretogranin II (SCG2). The non-target mRNA glutamate-cysteine ligase, catalytic subunit (GCLC) was used as a negative control.

To confirm the requirement for Star-PAP in the expression of these mRNAs, HEK 293 cells were treated with siRNA oligonucleotides specific for Star-PAP. Knockdown of Star-PAP expression resulted in a 5–15-fold decrease in the mRNA species examined (Fig. 5, *A* and *D*), indicating that Star-PAP is indeed required for their expression. Treatment of HEK 293 cells with CKI $\alpha$ -specific siRNA resulted in a dramatic decrease in HO-1 and NQO1 mRNAs, whereas other Star-PAP target mRNAs were unaffected (Fig. 5, *B* and *E*). Likewise, we found that treatment of cells with PIPKI $\alpha$ -specific siRNA resulted in comparable decreases in the same Star-PAP target mRNAs as CKI $\alpha$  siRNA, *viz.* HO-1 and NQO1 (Fig. 5, *C* and *F*). Together, these data raise the possibility that PIPKI $\alpha$  and CKI $\alpha$  may be working together to regulate specific Star-PAP target mRNAs.

HO-1 and NQO1 mRNAs encode important detoxifying enzymes involved in protection from reactive oxygen species and cellular injury (22). Both HO-1 and NQO1 are up-regulated in response to oxidative stress through increased transcription (23). We have shown previously that Star-PAP and



**FIGURE 3. CKI $\alpha$  can directly phosphorylate Star-PAP within its proline-rich region.** *A*, FLAG-tagged wild-type (wt) or K46R (kinase-dead (kd)) CKI $\alpha$  expressed in HEK 293 cells was purified and used to phosphorylate Star-PAP from the heat-inactivated FLAG-purified Star-PAP complex in an *in vitro* kinase assay. *B*, the addition of 50  $\mu$ M IC261 or PI-4,5-P<sub>2</sub> can block CKI $\alpha$  phosphorylation of Star-PAP. *C*, a schematic diagram depicts the Star-PAP truncations used. FL, full length; RRM, RNA recognition motif; NLS, nuclear localization signal; ZnF, zinc finger;  $\Delta$ ZF, zinc finger deletion. *D*, FLAG-Star-PAP was expressed in HEK 293 cells, purified by immunoprecipitation with anti-FLAG M2 antibody, and heat-inactivated prior to being subjected to *in vitro* phosphorylation by purified CKI $\alpha$  as described above. *E*, an alignment of the CKI $\alpha$  phosphorylation region in Star-PAP (amino acids 223–275) showing sequence conservation between mammalian species is shown. Serine and threonine residues are denoted with asterisks, and consensus CKI $\alpha$  sites are boxed.

PIPKI $\alpha$  are required for the up-regulation of the HO-1 transcript in response to a tBHQ-induced antioxidant response by playing a direct role in the 3'-end formation of this mRNA (6). Pretreatment of cells with the CKI-specific inhibitors CKI-7 and IC261 not only reduced the basal levels of HO-1 mRNA but also effectively blocked HO-1 induction after exposure to 100  $\mu$ M tBHQ (Fig. 5E). Treatment of HEK 293 cells with CKI $\alpha$ -specific siRNA oligonucleotides also caused a reduction in basal HO-1 levels; however, it did not block

HO-1 induction by tBHQ (Fig. 5F). The requirement for CKI $\alpha$  and PIPKI $\alpha$  specifically for Star-PAP target mRNAs involved in cellular protection suggests that phosphoinositide-based signaling is involved in the Star-PAP regulation of stress response mRNA maturation. However, although CKI inhibitors can effectively block HO-1 induction, CKI $\alpha$ -specific siRNA does not. This suggests that other CKI isoforms, or other protein kinases sensitive to CKI inhibitors, are also involved in the induction of HO-1 mRNA.

*CKI $\alpha$  Associates Specifically with Star-PAP-dependent mRNAs*—Although both Star-PAP and CKI $\alpha$  are required for the expression of the same mRNAs, it remains to be determined whether these proteins are actually acting together to regulate the levels of these mRNAs. Previously, we have shown that Star-PAP specifically interacts with its target mRNAs (6). Therefore, the ability of CKI $\alpha$  to interact with target mRNAs was examined by RNA immunoprecipitation (24).

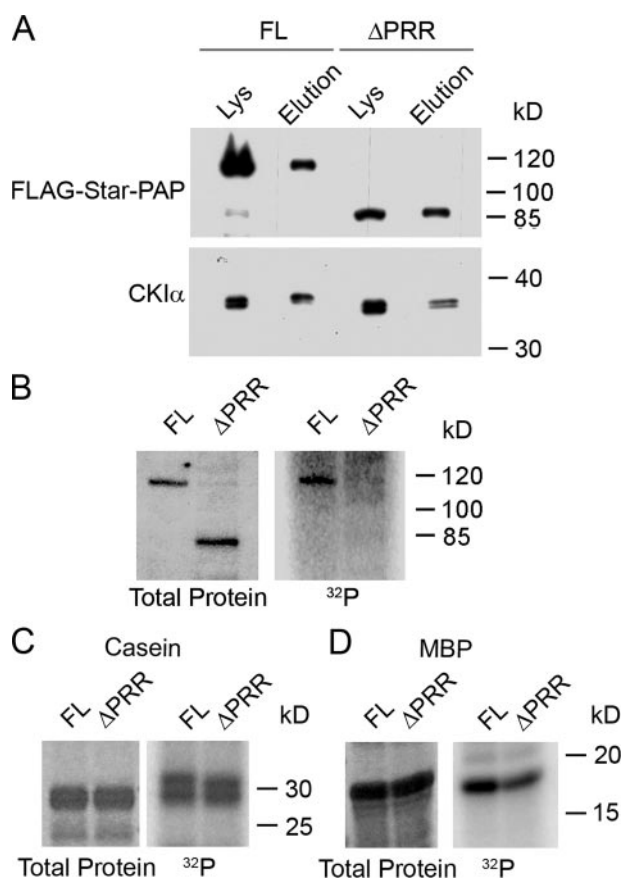
Endogenous CKI $\alpha$  and Star-PAP were immunopurified from HEK 293 cells, and total RNA was isolated from the immunoprecipitates. Specific mRNAs were then detected using reverse transcription-PCR. Similarly to Star-PAP, CKI $\alpha$  was specifically associated with its putative target mRNA, HO-1 (Fig. 6).

Notably, CKI $\alpha$  did not interact with the Star-PAP target mRNA CHAC1, the expression of which does not require CKI $\alpha$  or PIPKI $\alpha$ . This indicates that the association of CKI $\alpha$  with the Star-PAP complex occurs only with specific target mRNAs and that CKI $\alpha$  is not a universal component of all Star-PAP complexes.

## DISCUSSION

Star-PAP is a novel poly(A) polymerase involved in the regulation of specific mRNA expression levels. Star-PAP differs from both the canonical and non-canonical poly(A) polymerases in several ways, including the presence of the PRR insert in its catalytic domain, a unique association with the PI-4,5-P<sub>2</sub>-generating enzyme PIPKI $\alpha$ , and direct regulation by PI-4,5-P<sub>2</sub>. Here, we show that Star-PAP also differs markedly from canonical

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**FIGURE 4. Kinase activities and CKI $\alpha$  remain associated with Star-PAP when the proline-rich region is deleted.** *A*, full-length FLAG-Star-PAP (FL) and FLAG-Star-PAP $\Delta$ PRR complexes were expressed and purified from HEK 293 cells. The cell lysate (Lys) and the eluted FLAG affinity-purified complex are shown. Purified complexes were separated by SDS-PAGE and immunoblotted with anti-FLAG and anti-CKI $\alpha$  antibodies. Purified full-length FLAG-Star-PAP and FLAG-Star-PAP $\Delta$ PRR complexes were tested for associated kinase activity toward themselves (*B*), 100  $\mu$ g/ml casein (*C*), or 100  $\mu$ g/ml MBP (*D*) using *in vitro* protein kinase assays.

PAP $\alpha$  because of the association of protein kinase activity with the purified enzyme complex and that the PI-4,5-P<sub>2</sub>-sensitive protein kinase CKI $\alpha$  is one of the kinases contributing to this activity. We also found that both PIPKI $\alpha$  and CKI $\alpha$  are required for the synthesis of the Star-PAP target mRNAs HO-1 and NQO1, which are vital protective enzymes that respond to reactive oxygen species and cellular oxidative stress (22). Furthermore, like Star-PAP, CKI $\alpha$  is associated specifically with these mRNAs, suggesting that Star-PAP and CKI $\alpha$  work together to regulate their expression.

Phosphoinositide signaling is based on the subcellular targeting of phosphoinositide-generating enzymes and the subsequent production of phosphoinositide signaling molecules at specific subcellular sites (12). The presence of multiple PI-4,5-P<sub>2</sub>-sensitive components, including Star-PAP and CKI $\alpha$ , as well as the PI-4,5-P<sub>2</sub>-generating enzyme PIPKI $\alpha$ , suggests that the Star-PAP complex is a focal point for phosphoinositide signaling in the nucleus. Localized PI-4,5-P<sub>2</sub> generation by PIPKI $\alpha$  is likely able to modulate the activities of both CKI $\alpha$  and Star-PAP itself, which could result in the regulation of Star-PAP function by phosphoinositides at multiple levels.

It is interesting to note that the Star-PAP complex contains a number of proteins that are phosphorylated in this assay, none

of which are present in the PAP $\alpha$  complex (Fig. 1A). As with Star-PAP, phosphorylation of these proteins is also inhibited by PI-4,5-P<sub>2</sub>. This suggests that additional components of the Star-PAP complex may also be regulated by PI-4,5-P<sub>2</sub>-sensitive protein phosphorylation.

Combined, these data point to the Star-PAP complex as a site where multiple PI-4,5-P<sub>2</sub>-sensitive components are present to take advantage of locally generated PI-4,5-P<sub>2</sub> to regulate Star-PAP function. The presence of phosphoinositides in the nucleus has been well established; however, until recently, the mechanisms by which they affected nuclear events were unknown (25). The data presented here help to establish the mechanism by which nuclear phosphoinositides generated at nuclear speckles regulate mRNA processing and expression.

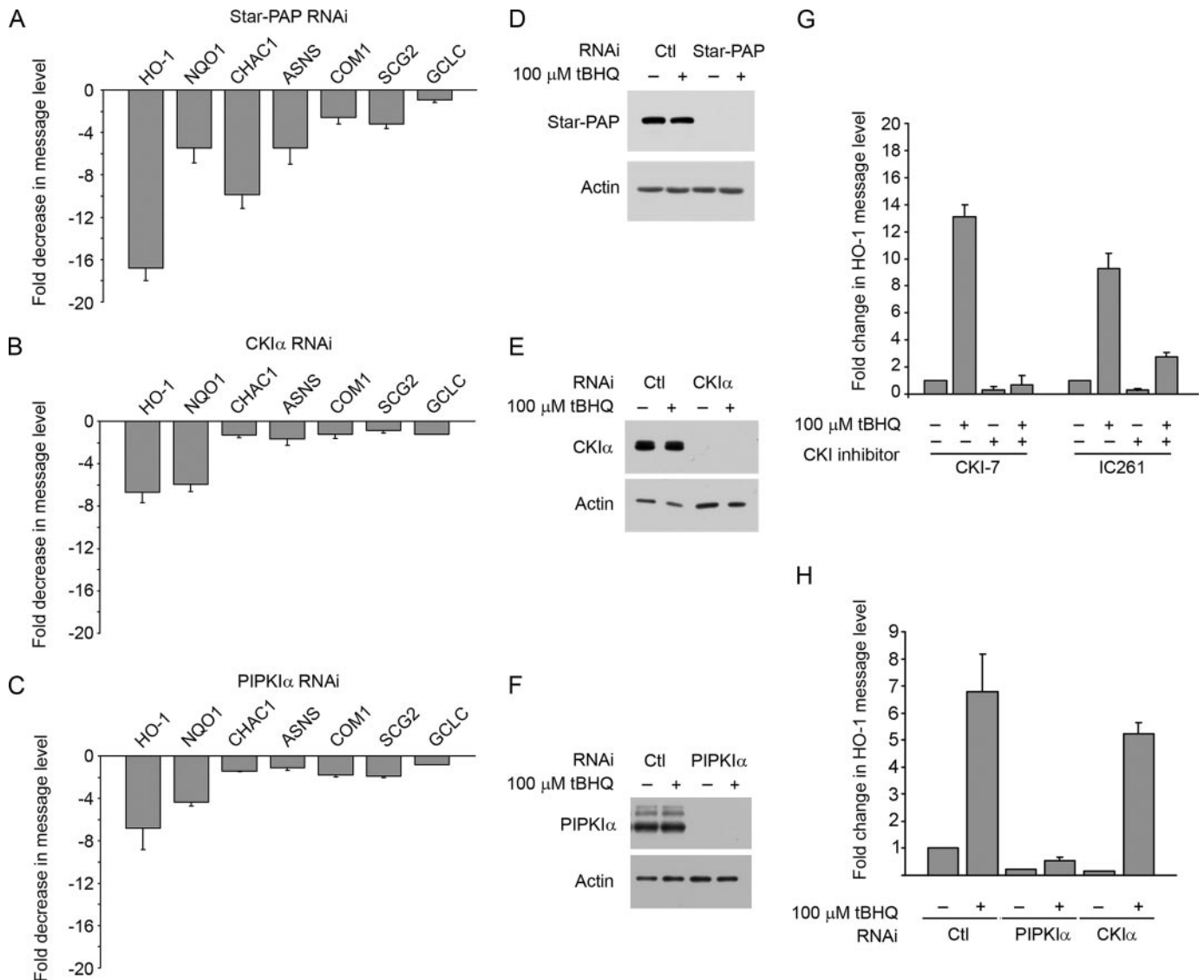
CKI $\alpha$  and PIPKI $\alpha$  are required for the expression of some but not all Star-PAP target mRNAs. This suggests that phosphoinositide-based signal transduction does not regulate all aspects of Star-PAP function and, additionally, that maturation of all Star-PAP target mRNAs is not regulated through the same signaling pathway. Furthermore, the fact that the related Star-PAP target mRNAs HO-1 and NQO1 require CKI $\alpha$  and PIPKI $\alpha$ , whereas unrelated Star-PAP target mRNAs such as CHAC1 do not, reveals that maturation of Star-PAP target mRNAs may be divided into groups based on their function and regulation. Star-PAP target mRNAs may exist as distinct groups, or modules, each of which is regulated by a different signal transduction pathway. For example, the HO-1/NQO1 oxidative stress response module is regulated by a CKI $\alpha$ - and PIPKI $\alpha$ -based phosphoinositide pathway, whereas other modules may be controlled by different pathways. This arrangement would allow Star-PAP to specifically affect the expression of individual mRNA groups.

It is also notable that CKI $\alpha$  is associated only with Star-PAP target mRNAs, the levels of which it affects. This indicates that the composition of the Star-PAP 3'-processing complex is dynamic and can change to regulate different groups of mRNAs. It is therefore likely that other, as yet unidentified signal transduction components can also be associated in a Star-PAP complex. This suggests a model in which different signal transduction components assemble into the Star-PAP complex to control the expression of specific Star-PAP target mRNA modules. Each of these different Star-PAP complexes would be targeted to a specific Star-PAP module. The activation of specific signal transduction pathways could then direct Star-PAP to participate in the 3'-end formation of select mRNAs to which that specific complex was targeted while not affecting the expression of the others. In this manner, Star-PAP would be positioned to interpret an array of stimuli and coordinate the cellular response by regulating mRNA expression.

It is still unclear how the Star-PAP complex identifies different mRNA modules. It could be that incorporation of certain factors directs the complex to different mRNA targets. Alternatively, association of the Star-PAP complex with members of each mRNA group could recruit different sets of signal transduction components. Identification of more Star-PAP target mRNAs and their inherent modules will be necessary before the mechanism of Star-PAP specificity can be explored.

Although basal expression of HO-1 requires both PIPKI $\alpha$  and CKI $\alpha$ , CKI $\alpha$  is not required for HO-1 induction, suggesting





**FIGURE 5. CKI $\alpha$  and PIPK1 $\alpha$  are required for the maintenance of specific Star-PAP mRNAs.** A–C, quantitative real-time PCR analysis of mRNA expression levels after treatment with siRNA oligonucleotides specific for Star-PAP (A), CKI $\alpha$  (B), or PIPK1 $\alpha$  (C) relative to treatment with a control (Ctl) siRNA oligonucleotide. D–F, immunoblotting of representative protein levels from cells used in A–C with anti-Star-PAP (D), anti-CKI $\alpha$  (E), or anti-PIPK1 $\alpha$  (F) antibodies to demonstrate the extent of siRNA knockdown. G, quantitative real-time PCR analysis of HO-1 message levels from cells treated with 100  $\mu$ M tBHQ after a 2.5-h pretreatment with the CKI inhibitor IC261 (50  $\mu$ M) or CKI-7 (250  $\mu$ M). H, quantitative real-time PCR analysis of HO-1 message levels from CKI $\alpha$  or PIPK1 $\alpha$  knockdown cells treated with 100  $\mu$ M tBHQ or Me<sub>2</sub>SO (control) for 4 h. Quantitative real-time PCR results are the average of three independent experiments. Error bars represent 1 S.D. RNAi, RNA interference. ASNS, asparagine synthetase; GCLC, glutamate-cysteine ligase catalytic subunit.

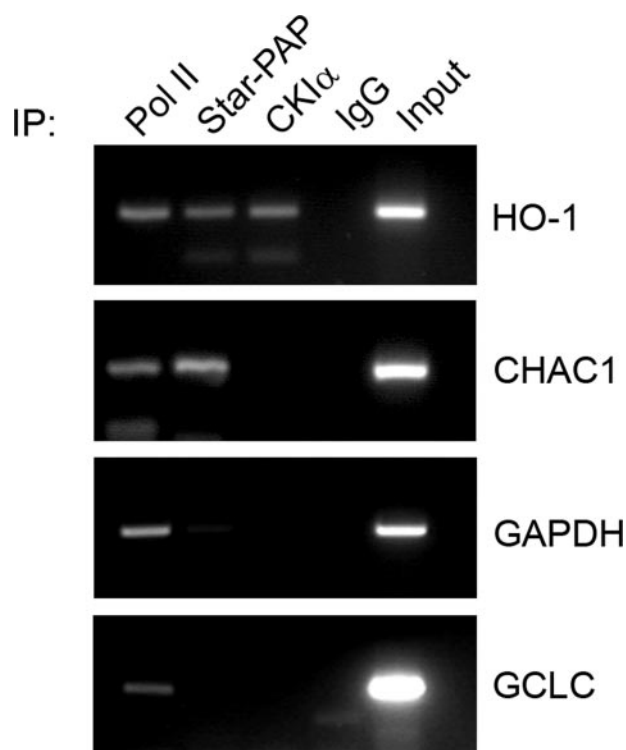
overlapping but unique roles for each protein in Star-PAP regulation. However, CKI inhibitors are effective in blocking tBHQ-induced HO-1 mRNA expression. This raises the possibility that other CKI isoforms can compensate for the loss of CKI $\alpha$  and can allow Star-PAP-dependent induction of HO-1 mRNA expression. Alternatively, it may be that other CKI isoforms, or even other CKI inhibitor-sensitive protein kinases, are a functional component of the Star-PAP complex and play a role in Star-PAP-dependent HO-1 mRNA induction.

The location of the CKI $\alpha$  phosphorylation site in the PRR of Star-PAP that interrupts the catalytic domain means that phosphorylation of Star-PAP could affect its catalytic activity. This could be through either a direct effect of the phosphorylation on activity or the modulation of an interaction with yet unknown regulatory proteins. Alternatively, it could be that

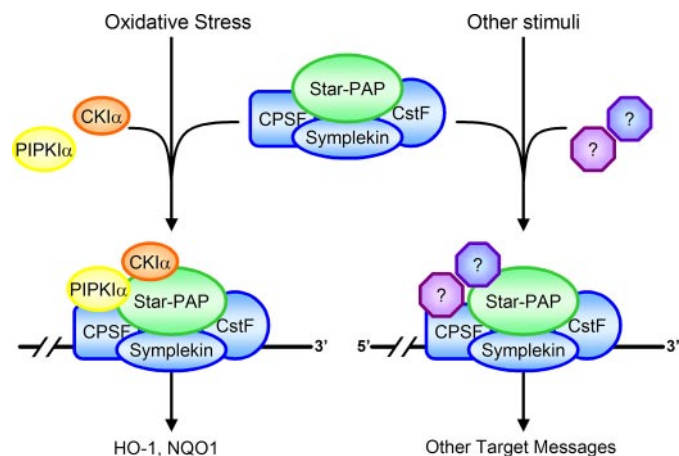
this site is somehow important for substrate recognition or even possibly involved in PI-4,5-P<sub>2</sub> binding and modulation of Star-PAP activity. Deletion of either the entire PRR or the portion that contains the CKI $\alpha$  phosphorylation site(s) results in a complete loss of Star-PAP poly(A) polymerase activity (data not shown). This may be due to a loss of regulatory elements, but just as likely, it may be due to a general disruption of the structure of the catalytic core of the Star-PAP enzyme.

Phosphorylation of PAP $\alpha$  has been shown to modulate its activity (26). However, on PAP $\alpha$ , this phosphorylation occurs on a serine/threonine-rich region in its carboxyl terminus. A comparable region does not exist in the same location on Star-PAP, so it is likely that the mechanism by which phosphorylation affects Star-PAP function will be different from that by which phosphorylation affects the activity of PAP $\alpha$ . We have

## Phosphorylation of Star-PAP by CKI $\alpha$



**FIGURE 6. CKI $\alpha$  specifically interacts with some Star-PAP target mRNAs.** RNA polymerase II (*Pol II*), Star-PAP, or CKI $\alpha$  was immunoprecipitated (*IP*) from nuclear extracts isolated from HEK 293 cells cross-linked with 1% formaldehyde. The cross-links were reversed, and total RNA was isolated from the immunoprecipitates and analyzed by reverse transcription-PCR with gene-specific primers for the Star-PAP targets HO-1 and CHAC1 as well as the non-targets glutamate-cysteine ligase catalytic subunit (*GCLC*) and glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*). A nonspecific rabbit IgG was used as a control.



**FIGURE 7. Model of Star-PAP complex assembly and target mRNA specificity.** Star-PAP exists as part of a basic 3'-processing complex that includes CPSF subunits, CstF subunits, and cleavage factors I and II. Signal transduction components are then integrated into a basal complex, forming a number of distinct Star-PAP complexes, each of which targets a different group of Star-PAP target messages. Each of these complexes is capable of 3'-mRNA processing and can respond to different signal transduction pathways. In this model, a CKI $\alpha$ /PIP1 $\alpha$  Star-PAP complex can regulate the expression of the oxidative stress response mRNAs HO-1 and NQO1.

not been able to identify the specific CKI $\alpha$  phosphorylation site on Star-PAP using individual point mutants. It is possible that CKI $\alpha$  is phosphorylating Star-PAP on multiple sites within the PRR. To gain a better understanding of how CKI $\alpha$  phosphorylation affects Star-PAP function, these site(s) must be identified.

Although CKI $\alpha$  is one protein kinase component of the Star-PAP complex, it is likely not the only one. The Star-PAP complex contains activity toward both MBP and casein. However, CKI $\alpha$  is unable to directly phosphorylate MBP, suggesting that there is at least one other protein kinase associated with the Star-PAP complex. Additionally, CKI inhibitors are capable of inhibiting casein phosphorylation by the associated kinase activity but are ineffective in blocking MBP phosphorylation (data not shown). Finally, most CKI $\alpha$  substrates must first be phosphorylated by another kinase to generate a CKI $\alpha$  phosphorylation site (20). Notably, CKI $\alpha$  is unable to phosphorylate *Escherichia coli*-expressed, purified Star-PAP (data not shown), which would not be expected to contain the priming phosphorylation.

Together, these data strongly suggest that another kinase is also associated with the purified Star-PAP complex. However, the contribution of this kinase to Star-PAP phosphorylation is still unclear. CKI inhibitors are able to block a majority of Star-PAP phosphorylation *in vitro*; however, the ability of this kinase to phosphorylate Star-PAP *in vivo* and/or afterward in response to specific stimuli cannot be discounted. It may be that this unidentified kinase activity represents the CKI $\alpha$  priming kinase. Alternatively, this kinase may be involved in the regulation of other Star-PAP mRNAs, consistent with our model in which distinct Star-PAP complexes are each responsible for controlling the expression of specific mRNA modules.

The data presented here demonstrate an unexpected new nuclear substrate and function for the PI-4,5-P<sub>2</sub>-sensitive protein kinase CKI $\alpha$ . CKI $\alpha$  can directly phosphorylate Star-PAP and is required for the expression of the Star-PAP target mRNAs HO-1 and NQO1, which encode critical detoxifying enzymes. The involvement of Star-PAP, PIP1 $\alpha$ , and CKI $\alpha$  in the expression of these mRNAs suggests that a nuclear phosphoinositide signaling pathway regulates the level of specific mRNAs in response to oxidative stresses. However, the finding that phosphoinositide-based regulation of Star-PAP affects only a subset of Star-PAP target mRNAs demonstrates that Star-PAP is a very sophisticated regulatory enzyme. The association of the basic Star-PAP 3'-processing complex with distinct signal transduction components allows Star-PAP to specifically regulate the expression of discrete groups of mRNAs. In this manner, as depicted in Fig. 7, Star-PAP is capable of interpreting signals from distinct pathways and translating them into unique mRNA expression profiles by modulating the 3'-end formation of specific groups of Star-PAP target mRNAs.

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