

Supplementary information to:

Unraveling the stepwise maturation of the yeast telomerase including a Cse1 and Mtr10 mediated quality control checkpoint

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Supplementary Tables

Supplementary Table 1. Yeast strains used in this study. Related to Figures 1-6

Number	Genotype	Source
HKY36	<i>MATa ura3-52 leu2Δ1 his3Δ200</i>	1
HKY37	<i>MATa ura3-52 leu2Δ1 his3Δ200 srp1-31</i>	2
HKY46	<i>MATa ura3-52 lys2-301 ade2 mtr10-1</i>	3
HKY208	<i>MATa ura3-52 ade2-101 his3-11,15, trp1-Δ901 cse1-1</i>	4
HKY316	<i>MATa ura3-52 leu2Δ1 trp1Δ63 MTR10-9xMyc-TRP1</i>	This study
HKY380	<i>MATa his3Δ1 leu2Δ0 met12Δ0 ura3Δ0 npl3::KanMX4</i>	Euroscarf
HKY644	<i>MATa ade2, his3, leu2, trp1, ura3 mex67::HIS3 pUN100-mex67-5 (LEU2, CEN)</i>	5
HKY1028	<i>MATa his3Δ1; leu2Δ0; lys2Δ0; ura3Δ0 rrp6::kanMX4</i>	Euroscarf
HKY1073	<i>MATa his3Δ1; leu2Δ0; met15Δ0; ura3Δ0; yKu70::kanMX4</i>	Euroscarf
HKY1079	<i>MATa his3Δ1; leu2Δ0; met15Δ0; ura3Δ0; RAP1-GFP::HIS3MX6</i>	7
HKY1093	<i>MATa his3Δ1; leu2Δ0; met15Δ0; ura3Δ0; CDC13-GFP::HIS3MX6</i>	7
HKY1193	<i>Tgs1::KanMX4/Tgs1::KanMX4 his3Δ1 / his3Δ1; leu2Δ0 / leu2Δ0; ura3Δ0 / ura3Δ0; lys2Δ0 / LYS2; MET15 / met15Δ0</i>	Euroscarf
HKY1293	<i>MATa ura3-52 lys2-801 trp-Δ1 his3-Δ200 leu2-Δ1 tlc1- Δ::HIS</i>	6
HKY1277	<i>MATa his3Δ1 leu2Δ0 met15Δ0 ura3Δ0 MTR10- GFP::HIS3MX6</i>	7
HKY1353	<i>MATa ura3-52 mex67::HIS3 xpo1::TRP1 pUN100 (CEN LEU2) mex67-5 xpo1-1::HIS3</i>	8
HKY1596	<i>MATa his3Δ1 leu2Δ0 met15Δ0 ura3Δ0 CSE1- GFP::HIS3MX6</i>	7
HKY1642	<i>MATa ura3-52 lys2-801 ade2-Δ trp1-Δ63 his3-Δ200 leu2- Δ1 smb::KAN pGal-TRP1-SmB</i>	9
HKY1645 (<i>smb smd1</i>)	<i>MATa ura3-52 lys2-801 ade2-Δ trp1-Δ63 his3-Δ200 leu2- Δ1 smb::KAN smd1::LEU2</i>	9
HKY1689	<i>MATa rrp6::kanMX4 mex67::HIS3 pUN100-mex67-5 (LEU2, CEN)</i>	10
HKY1776	<i>MATa mtr10:kanMX4 lys ura leu his</i>	This study
HKY1799	<i>MATa his3Δ1 leu2Δ0 met15Δ0 ura3Δ0 CBP20- GFP::HISMX6</i>	Euroscarf
HKY1815	<i>MATa ura leu Pop1-GFP::HisMX6</i>	Euroscarf
HKY2087	<i>MATa his ura KAN cse1-1 mtr10:kanMX4</i>	This study
HKY2093	<i>MATa ura cse1-1 Pop1-GFP::HisMX6</i>	This study
HKY2101	<i>MATa leu2Δ1 ura3-52 MTR10-9xMyc-TRP1 Pop1- GFP::HisMX6</i>	This study
HKY2153	<i>MATa ura Pop1-GFP::HisMX6 mex67::HIS3 pUN100-mex67-5 (LEU2, CEN)</i>	This study
HKY2259	<i>MATa; ; ura3-52, ade2-101, trp1-Δ901cse1- RAP1-GFP::HIS3MX6</i>	This study
HKY2261	<i>MATa; ; ura3-52, ade2-101, trp1-Δ901cse1- CDC13-GFP::HIS3MX6</i>	This study

Supplementary Table 2. Plasmids used in this study. Related to Figures 1-6.

Number	Genotype	Name
pHK88	CEN URA3	11
pHK206	CEN URA CSE1	This study
pHK765	CEN URA GFP-Npl3	12
pHK1469	CEN URA SmB-GFP	13
pHK1483	CEN URA GFP-POP1	14
pHK1589	URA3 EST1-(Gly)6-(myc)12	15
pHK1606	CEN URA pAdh-Est1-GFP	This study

Supplementary Table 3. Oligonucleotides used in this study. Related to Figures 1-4, 6. Forward primer (fw) and reverse primer (rev).

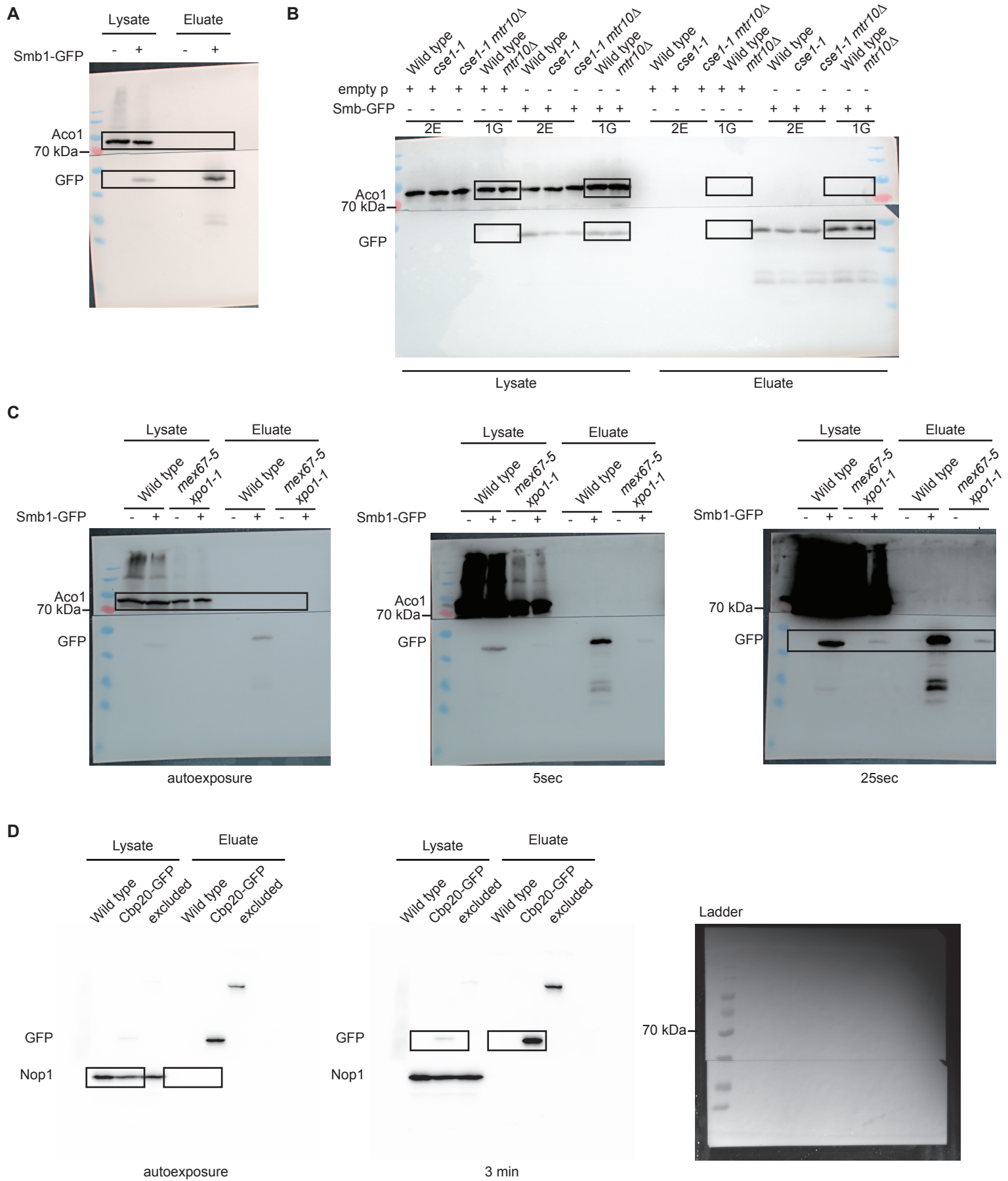
Number	Sequence	Name
HK1002	5'-TGCTAAGGCTGTCCGTAAGG-3'	<i>Tdh1</i> fw
HK1003	5'-TCAGAGGAGACAACGGCATC-3'	<i>Tdh1</i> rev
HK1384	5'-GCGGAAGGAACCGTGTGTTC-3'	<i>TLC1</i> immature fw
HK1385	5'-GAAGCCTACCATCACCACACC-3'	Internal <i>TLC1</i> fw
HK1386	5'-ACAGCGCTTAGCACCGTCTG-3'	Internal <i>TLC1</i> rev
HK1539	5'-DIG-CCACCACACACACCCACACCC-3'	5' DIG labeled CA – Telomeric repeat probe
HK1598	5'-GGCCCCAGGTAAGAAAGTCG-3'	RPL8 fw
HK1599	5'-GAAGGTTTCGGCAGCGGTG-3'	RPL8 rev
HK1738	5'-TGCAAACCTCCTTGGTCACAC-3'	U1 snRNA (snR19) fw
HK1739	5'-CCAGGCAGAAGAAACAAAGG-3'	U1 snRNA (snR19) rev
HK1761	5'-CY3- GCGCACACACAAGCATCTACTGACACCAGCAT ACTCGAAATTCTTTGG-CY3-3'	<i>TLC1</i> probe 1
HK1789	5'-CY3- CGATAAGATAGACATAAAGTGACAGCGCTTAGCA CCGTCTGTTGC-CY3-3'	<i>TLC1</i> probe 2
HK1790	5'-CY3- CCTACTCGTATTTTTCTCTGTCACATCGTTCGATGT ACGGGGCACATTTGG-CY3-5'	<i>TLC1</i> probe 3
HK2154	5'-CCAGAACAATCCGTACACAAGG-3'	<i>Hem15</i> fw
HK2155	5'-GCAATTGTCTTCTGATACTTAGCAC-3'	<i>Hem15</i> rev
HK2859	5'-CAGCTTTACAGATCAATGGC-3'	U5 snRNA (snR7-L) fw
HK2860	5'-TATGGCAAGCCCACAGTAA-3'	U5 snRNA (snR7-L) rev
HK3089	5'-AGTTACGCTAGGGATAACAGGG-3'	21S rRNA fw
HK3090	5'-TGACGAACAGTCAAACCCTTC-3'	21S rRNA rev
HK3513	5'-ACGCGCGATTCTACAATAC-3'	<i>TLC1</i> immature rev

Supplementary references:

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14. Gill T, Aulds J, Schmitt ME. A specialized processing body that is temporally and asymmetrically regulated during the cell cycle in *Saccharomyces cerevisiae*. *The Journal of cell biology* **173**, 35-45 (2006).
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Supplementary Fig. 1



Supplementary Figure 1. Original western blots of Figure 1. The PageRuler Prestained (Thermo Scientific #26616) was used as Ladder.

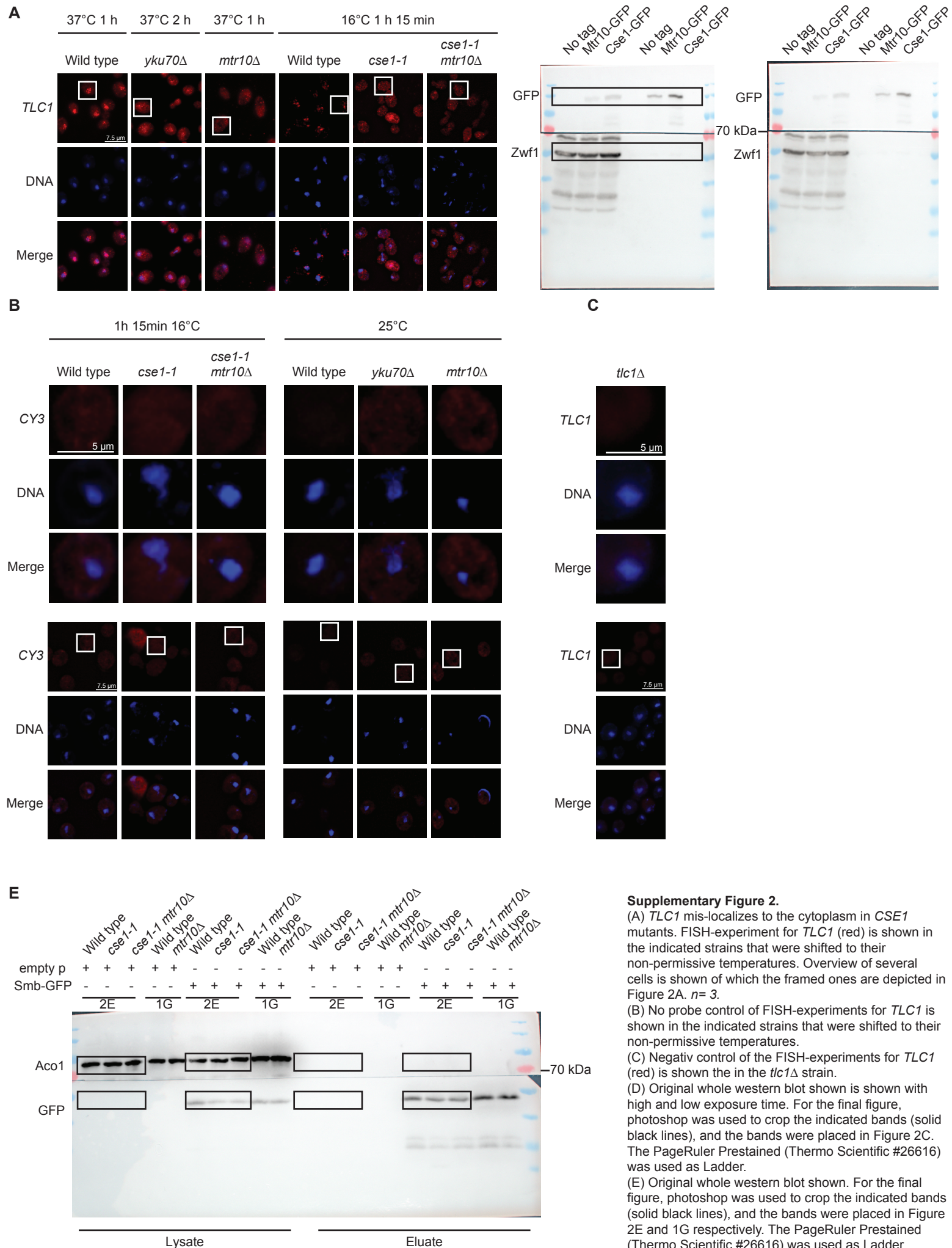
(A) Original whole western blot shown. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Figure 1A.

(B) Original whole western blot shown. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Figure 1G and 2E respectively.

(C) Original whole western blot with different exposure times is shown. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Figure 1C.

(D) Original whole western blot with different exposure times is shown. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Figure 1J.

Supplementary Fig. 2



Supplementary Figure 2.

(A) *TLC1* mis-localizes to the cytoplasm in *CSE1* mutants. FISH-experiment for *TLC1* (red) is shown in the indicated strains that were shifted to their non-permissive temperatures. Overview of several cells is shown of which the framed ones are depicted in Figure 2A. $n = 3$.

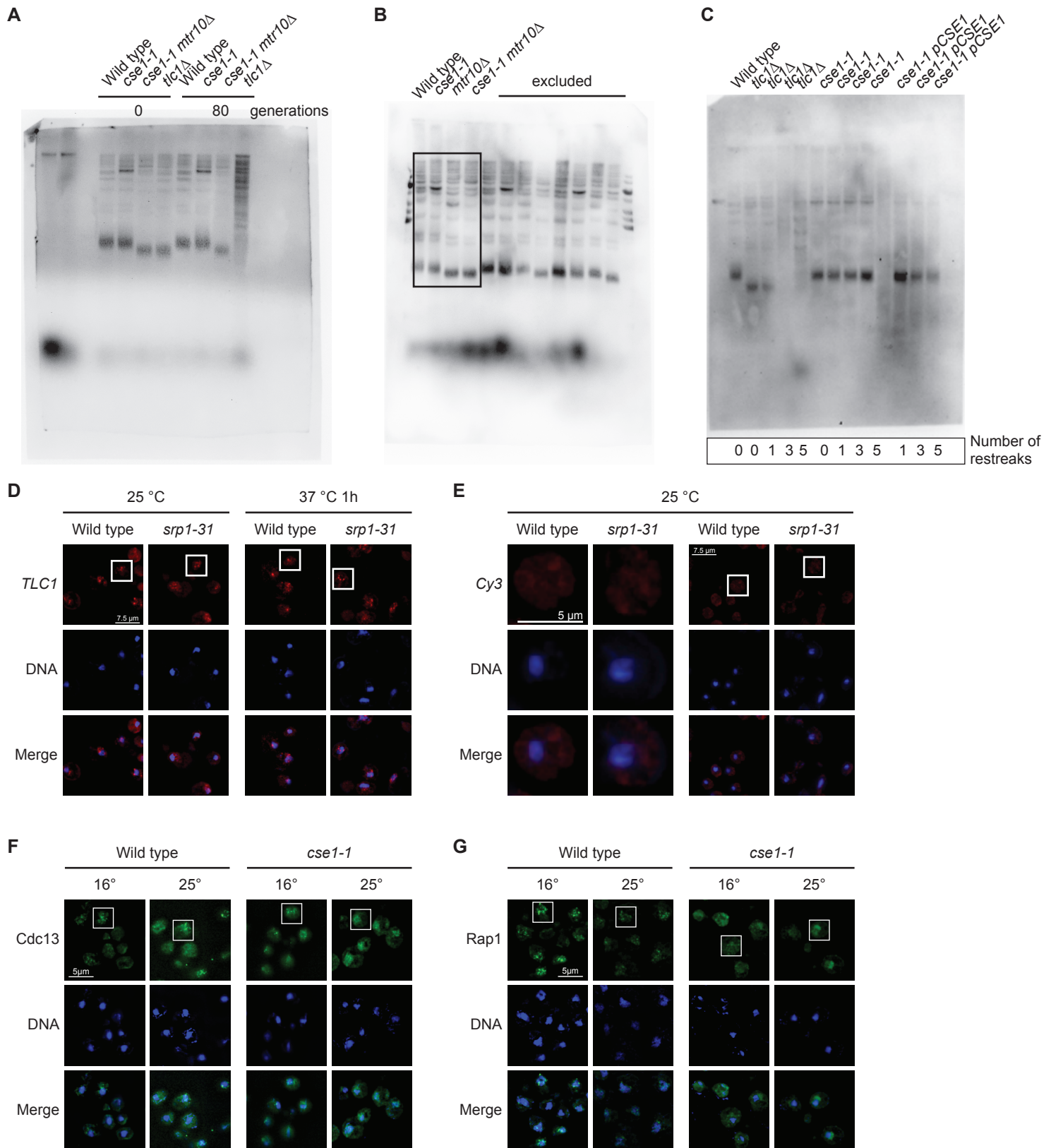
(B) No probe control of FISH-experiments for *TLC1* is shown in the indicated strains that were shifted to their non-permissive temperatures.

(C) Negative control of the FISH-experiments for *TLC1* (red) is shown in the *tlc1Δ* strain.

(D) Original whole western blot shown with high and low exposure time. For the final figure, photoshop was used to crop the indicated bands (solid black lines), and the bands were placed in Figure 2C. The PageRuler Prestained (Thermo Scientific #26616) was used as Ladder.

(E) Original whole western blot shown. For the final figure, photoshop was used to crop the indicated bands (solid black lines), and the bands were placed in Figure 2E and 1G respectively. The PageRuler Prestained (Thermo Scientific #26616) was used as Ladder.

Supplementary Fig. 3



Supplementary Figure 3.

(A) Whole southern blot is shown. XhoI digested genomic DNA of the indicated strains was used for the southern blot. The chromosome ends were detected with a digoxigenin labeled probe. Photoshop was used to adjust the levels so that the bands were more apparent. The bands were placed in Figure 3B.

(B) (A) Whole southern blot is shown. XhoI digested genomic DNA of the indicated strains was used for the southern blot. The chromosome ends were detected with a digoxigenin labeled probe. Photoshop was used to adjust the levels so that the bands were more apparent and the cropped area (solid black lines) is shown in Figure 3C.

(C) Whole southern blot is shown. XhoI digested genomic DNA of the indicated strains was used for the southern blot. The chromosome ends were detected with a digoxigenin labeled probe. Photoshop was used to adjust the levels so that the bands were more apparent. The bands were placed in Figure 3D.

(D) *TLC1* localization is not altered in the *srp1-31* mutant. FISH experiments show no mis-localization of *TLC1* in *srp1-31* after shift to 37 °C for 1h. Overview of several cells is shown of which the framed ones are depicted in (Fig. 3F) $n=3$.

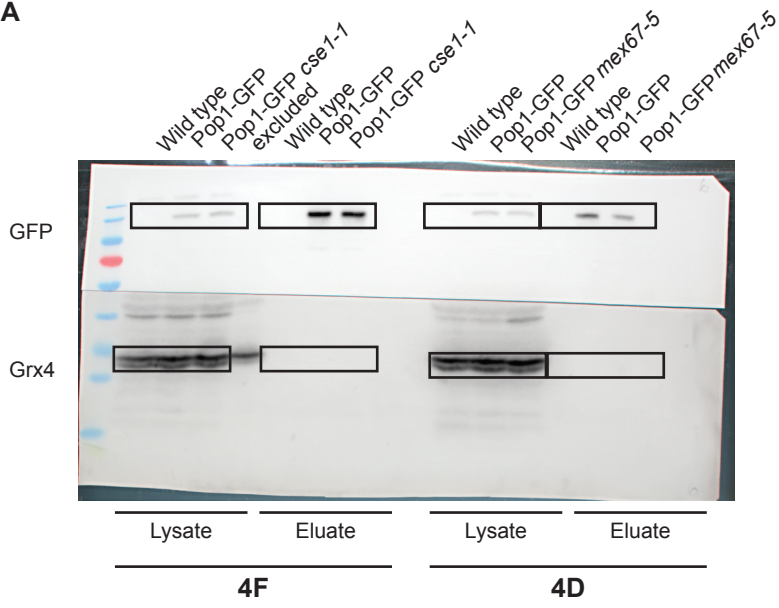
(E) No probe control of FISH-experiments for *TLC1* is shown in the *srp1-31* mutant.

(F) *Cdc13* localization is not altered in the *cse1-1* mutant. Immunofluorescence experiments with an antibody against GFP show no mis-localization of *Cdc13* in *cse1-1* after shifting the cells to 16 °C for 1h and 15min. Overview of several cells is shown of which the framed ones are depicted in (Fig. 3G) $n=3$.

(G) *Rap1* localization is altered in the *cse1-1* mutant. Immunofluorescence experiments with an antibody against GFP show nucleolar mis-localization of *Rap1* in *cse1-1* at the permissive temperature and a cytoplasmic mis-localization after shifting the strain to 16 °C for 1h and 15min. Overview of several cells is shown of which the framed ones are depicted in (Fig. 3H) $n=3$.

Supplementary Fig. 4

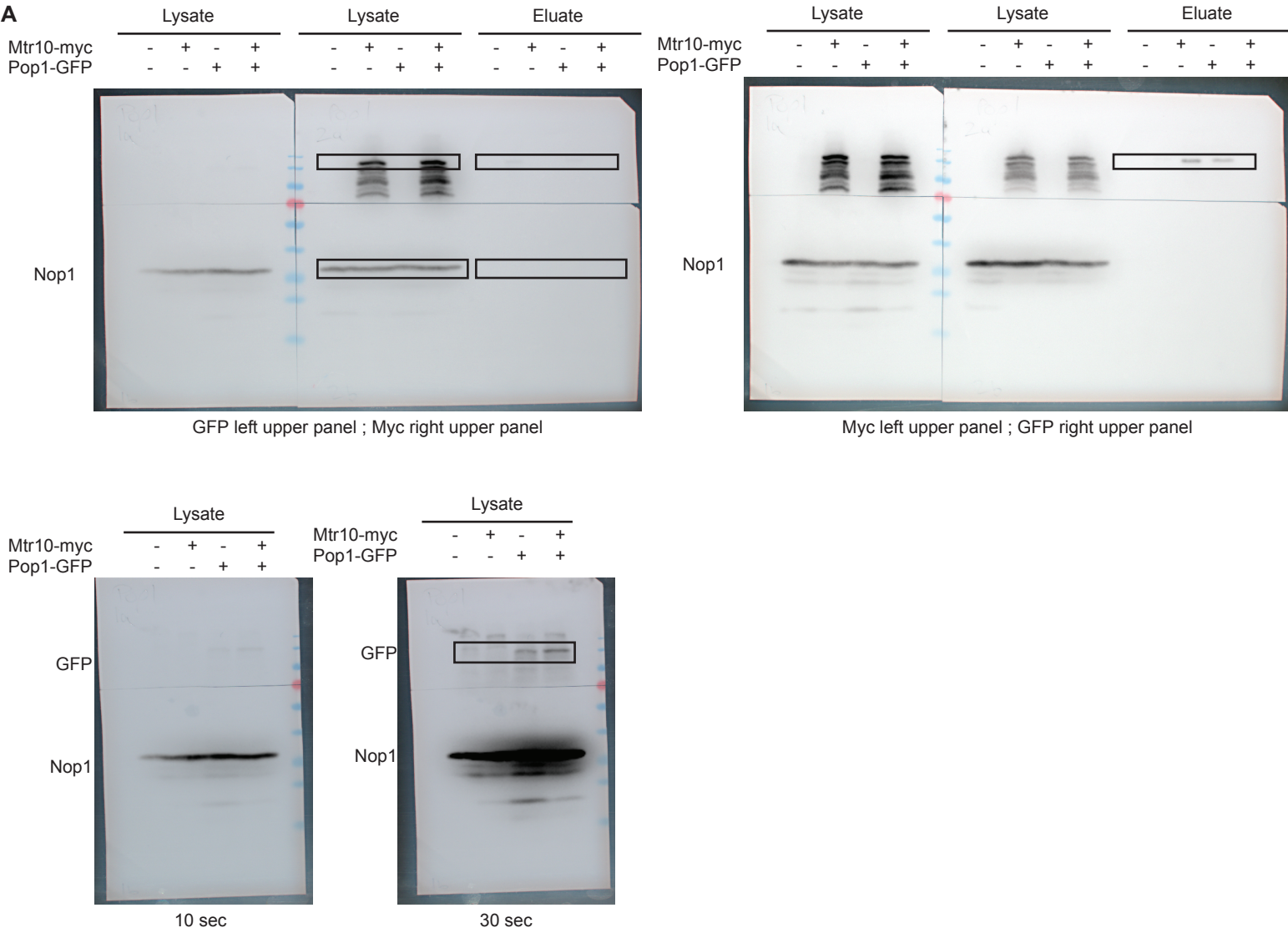
A



Supplementary Figure 4. Original western blots of Figure 4. The PageRuler Prestained (Thermo Scientific #26616) was used as Ladder

(A) Original whole western blot shown. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Fig. 4F and 4D respectively.

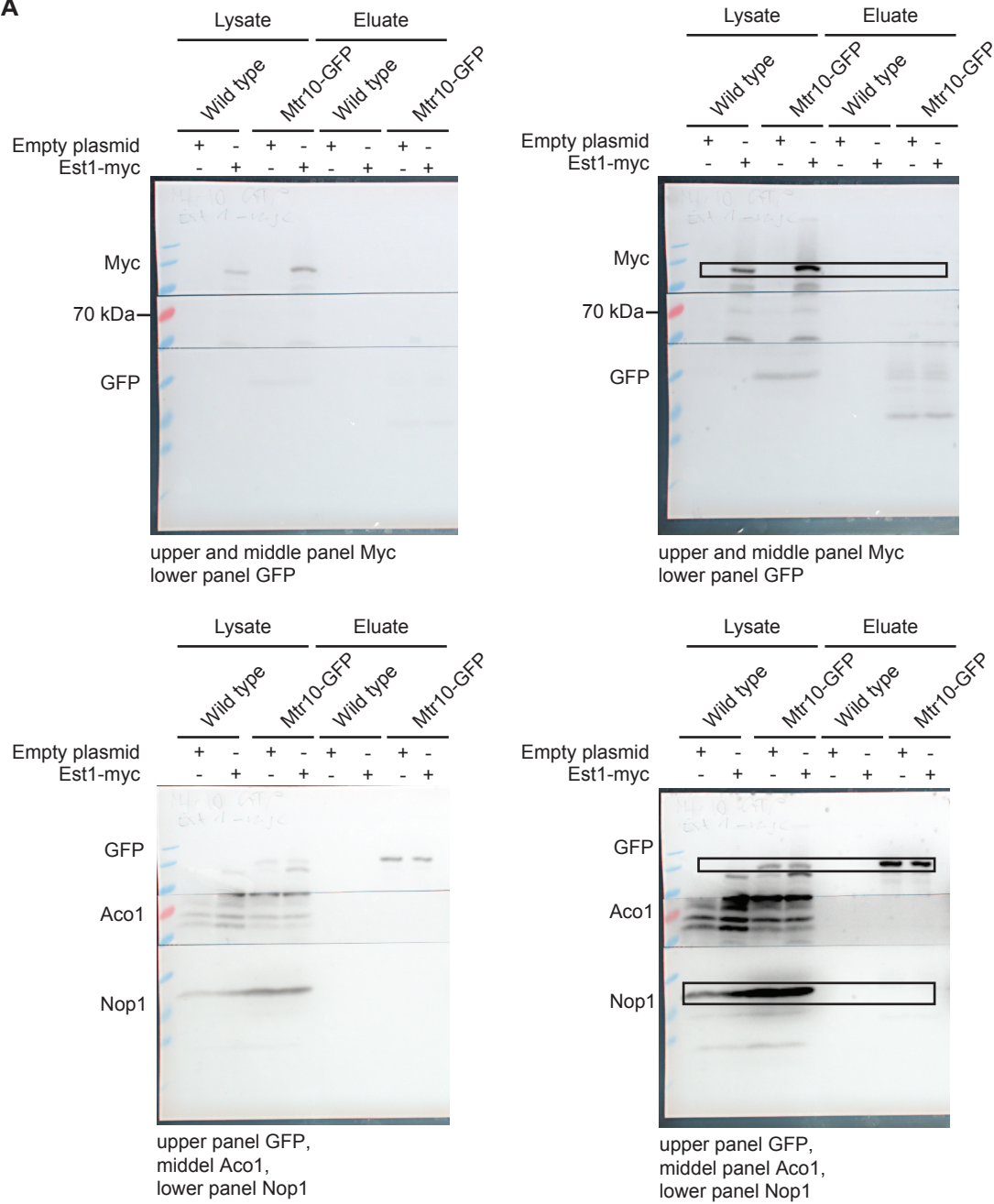
Supplementary Fig. 5.1



Supplementary Figure 5.1. Original western blots of Figure 5A.
The PageRuler Prestained (Thermo Scientific #26616) was used as Ladder
 (A) Original whole western blot shown. The lysates were loaded twice as the tagged proteins have almost identical size. Detection order is indicated underneath the blots. In the Eluate, the first step was to detect the protein to be co-precipitated. Afterwards, pull down detection was carried out. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Fig. 5A.

Supplementary Fig. 5.2

A



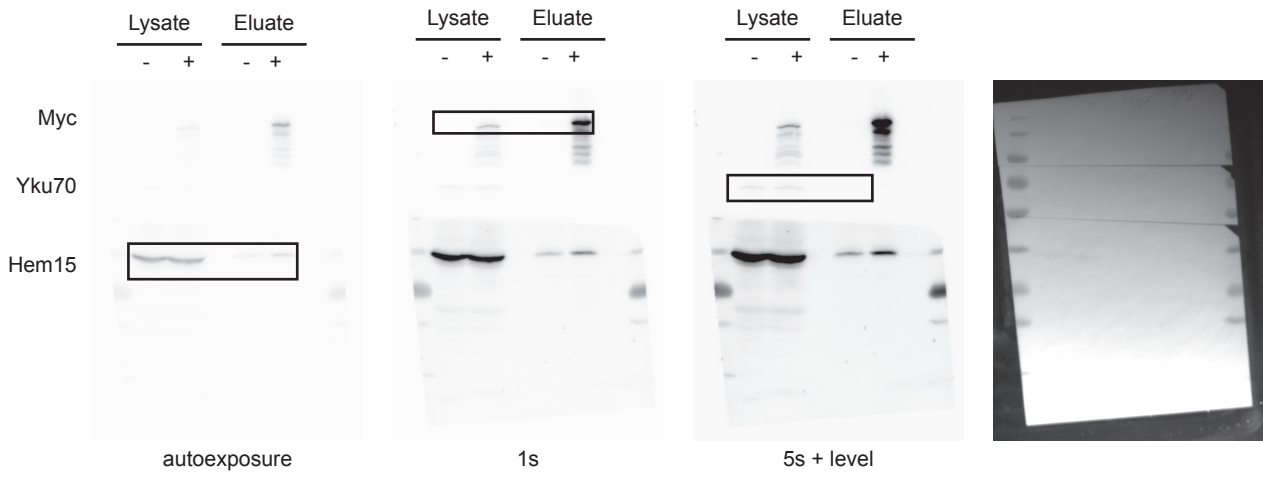
Supplementary Figure 5.2. Original western blots of Figure 5B.

The PageRuler Prestained (Thermo Scientific #26616) was used as Ladder

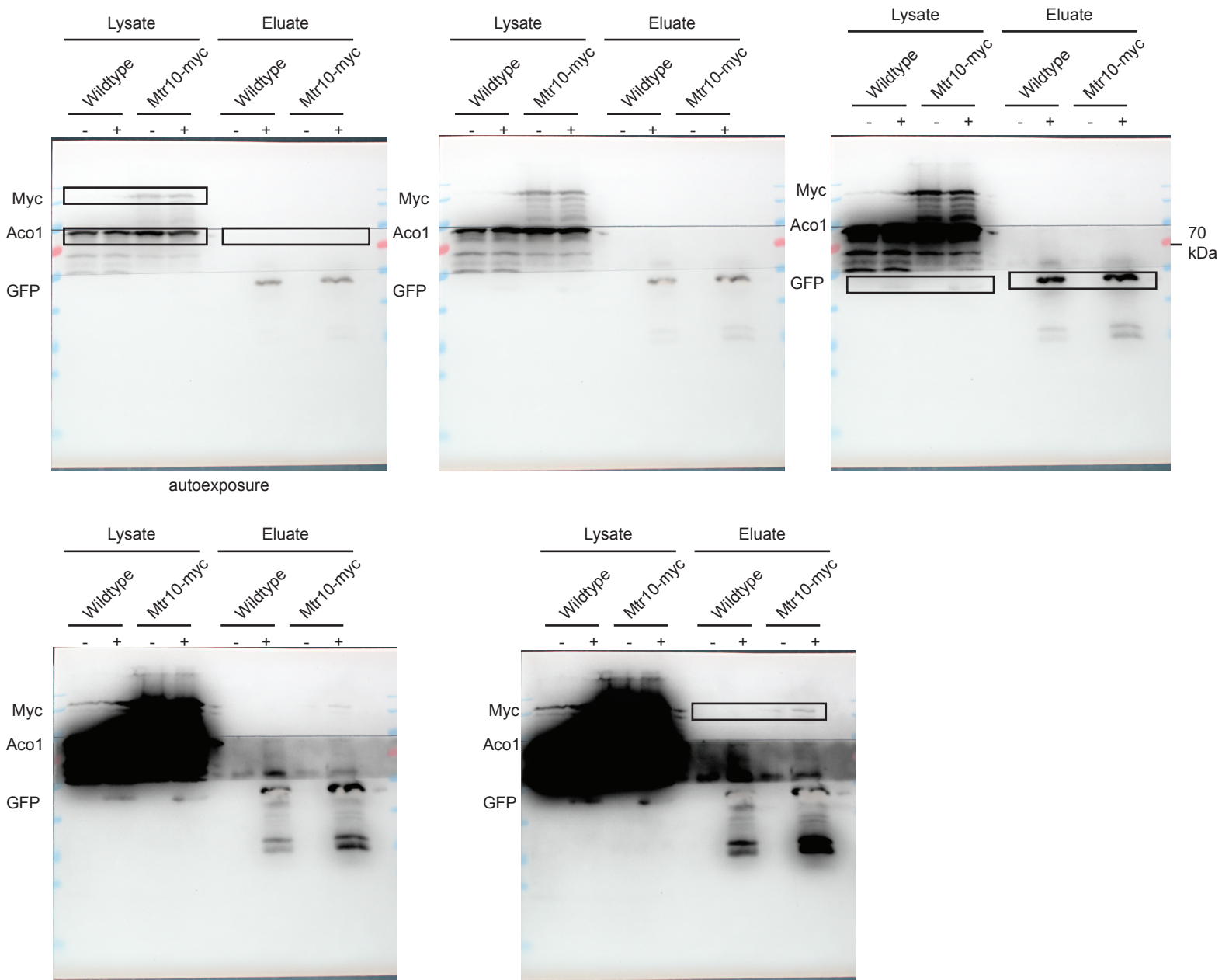
(A) Original whole western blot shown. Detection order is indicated underneath the blots. In the ELuate, the first step was to detect the protein to be co-precipitated. Afterwards, pull down detection was carried out. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Fig. 5B.

Supplementary Fig. 5.3

A



B



Supplementary Figure 5.3. Original western blots of Figure 5C and 5D.

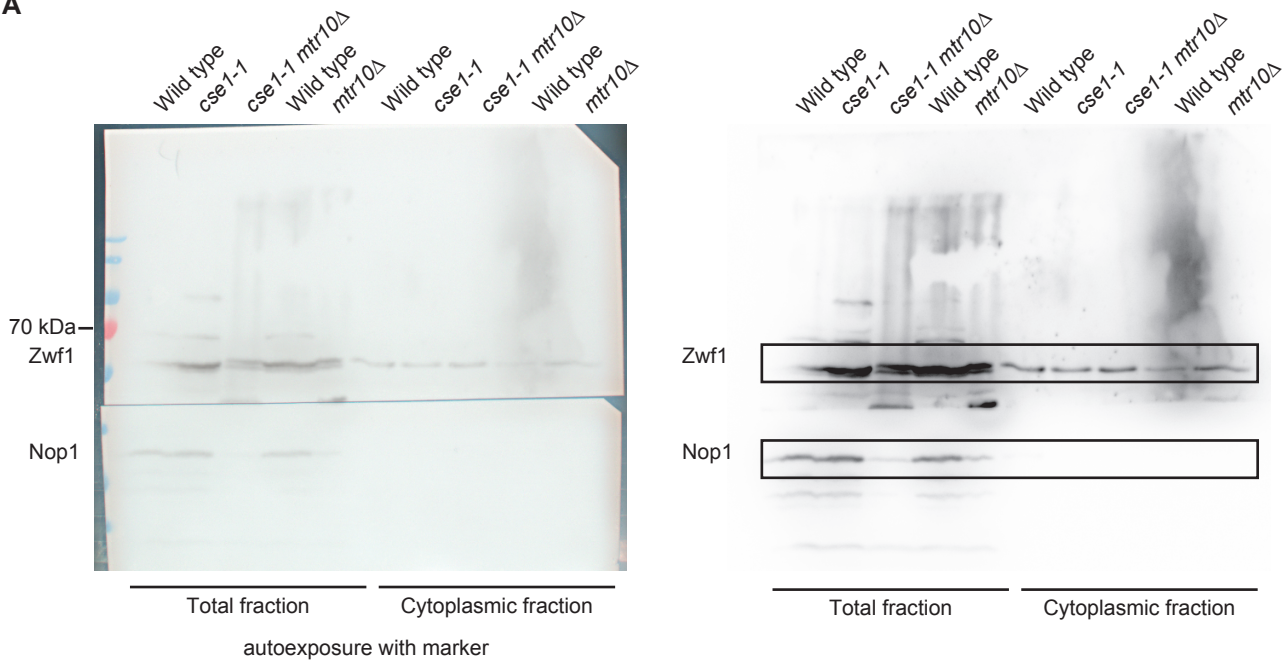
The PageRuler Prestained (Thermo Scientific #26616) was used as Ladder

(A) Original whole western blot is shown with different exposure times. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Fig. 5C.

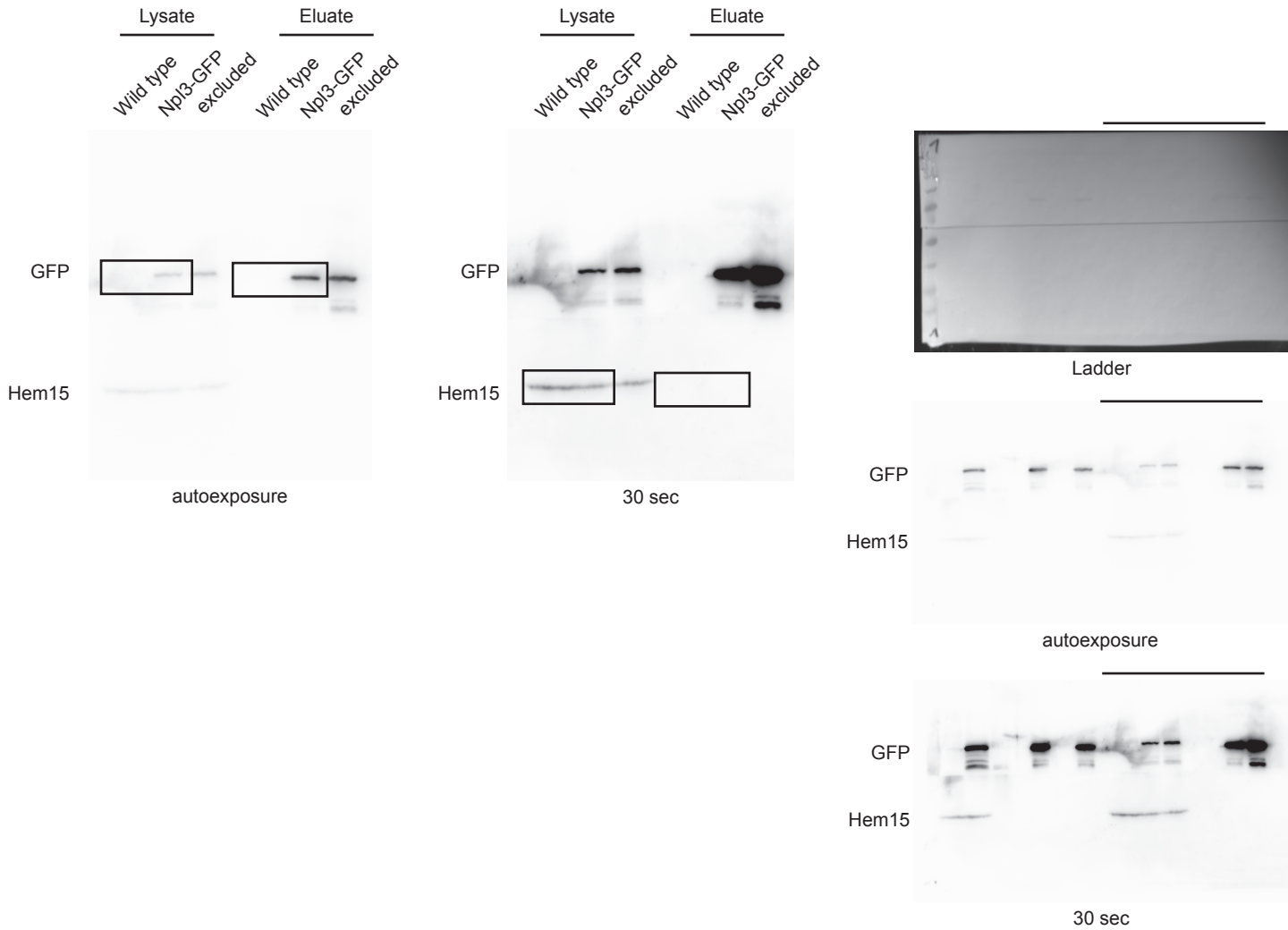
(B) Original whole western blot is shown with different exposure times. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Fig. 5D.

Supplementary Fig. 6

A



B



Supplementary Figure 6.

(A) Original whole western blot of a nucleo-cytoplasmic fractionation is shown. For the final figure, photoshop was used to enhance the levels and to crop the indicated bands (solid black lines) and the bands were placed in Figure 6A.
 (B) Original whole western blot of an RIP experiment is shown. For the final figure, photoshop was used to crop the indicated bands (solid black lines) and the bands were placed in Figure 6C.