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Supplemental Information

Global Analysis of Eukaryotic mRNA Degradation Reveals Xrn1-Dependent Buffering of Transcript Levels

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A


B


Figure S1 Expression level of genes in pat14 plotted against their position in chromosomes. Related to Figure 1
A. An aneuploid strain variant shows two-fold overexpression of genes on Chr II compared to the median level of all other genes.
B. The strain used in this study with normal ploidy. Such ploidy analysis was carried out for all strains used in this study. The analysis revealed that the published strain pop24 (Sun, M., et.al. 2012 Genome Research) has a chromosome IIX aneuploidy. A new knock-out strain was therefore generated by homologous recombination and measured by cDTA. The new data however does not change any of the conclusions in our previous work.


Figure S2 In vitro transcription assay. Related to Figure 2
A. The assay was carried out using nuclear extract from BY4741 (left) and xrn14 (right). TAPpurified Xrn1 and Xrn1pm has been added to the assay separately. RNase inhibitor RiboLock (Fermentas) was added to inhibit non-specific RNA degradation by enzymes present in the extract. The inhibitor is necessary to observe RNA synthesis and according to the manufacturer does not inhibit Xrn1. No difference was observed in the GCN4 activator dependent transcription activity in vitro from HIS4 promoter.
B. To check whether a nuclear extract prepared according to our protocol contains Xrn1 protein, we repeated extract preparation from a yeast strain carrying a TAP tag on Xrn1 that enabled protein monitoring in the absence of an antibody against yeast Xrn1. Western blotting revealed a specific protein degradation pattern that arises from the proteolytically sensitive Xrn1 protein (lane 4). A very similar pattern is observed in whole cell extracts (lane 1) but not in a control extract preparation from a strain that lacks the TAP tag (lane 2).
C. To test whether the recombinant Xrn1 protein we added to the assay was active as a nuclease, we incubated it with a 30 -mer RNA substrate with 5 ' monophosphate group (sequence: 5'-Phos/CGGCGGUCUUAUGUAACGGCGGUGUAAAUCUU-3') and observed partial degradation of the RNA by non-specific staining after 30-60 min of incubation (lanes 2, 3).


Figure S3 Scatter plot of cDTA analysis on $n r g 1 \Delta$ and $x r n 1 \Delta$ double knock out strain Scatter plot show the SR and DR changes in $n r g 1 \Delta x r n 1 \Delta$ double mutant (Related to Figure 3)


Figure S4 Growth analysis during cycloheximide perturbation. Related to Figure 4
$(A, B)$ Growth effect of cycloheximide on wild-type BY4741 cells. Cycloheximide was added at OD600 $=0.8$, and a concentration of $0-50 \mu \mathrm{~g} / \mathrm{ml}$ was tested.
(C) Replicates experiments with the concentration used in this study.


Figure S5. Cluster analysis provides evidence for a general mRNA degradation machinery.
The plot shows the color-coded median t-statistics of DRs of mRNAs that show significantly decreased or increased DRs in the gene deletion strains shown on the x -axis. The t -statistics gives the ratio of the difference in mean DR and its standard errors as a measure for differential stabilization (decreased/increased DR). Destabilization (increased DR) is shown in red, stabilization (decreased DR) in blue. The overall directionality of perturbed degradation indicates a general mRNA degradation machinery. (Related to Figure 6)


Figure S6 Antagonistic effect in $d c s 1$ and $d c s 2$. Related to Figure 6
Scatter plots show global changes in mRNA DRs (log fold, x-axis) vs. the global changes in SRs (log fold, y -axis). The global changes of $d c s 1$ and $d c s 2$ are plotted. The coordinate of the center of each circle is determined by the median DR and SR of the mutant. The diameters of the circles represent the relative comparison of the fold of RNA amount over wild-type level.

Table S1 Strains used in this study

| 1 | BY4743 | MAT $\boldsymbol{\alpha} / \mathbf{a}$; <br> his341/his341;leu240/leu240; <br> lys240/LYS2; MET15/met1540; <br> ura340/ura340 | OpenBiosystems |
| :---: | :---: | :---: | :---: |
| 2 | BY4741 | MAT a his341; leu240; met1540; ura340 | OpenBiosystems |
| 3 | air1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met $15 \Delta 0$; ura3 00 ; yil079c::kanMX4 | OpenBiosystems |
| 4 | air2 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 10 ; ura3 30 ; ydl175c::kanMX4 | OpenBiosystems |
| 5 | bud13 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 $\Delta 0$; ura3 $\Delta 0$; ygl174w::kanMX4 | OpenBiosystems |
| 6 | caf40 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 0 ; ura3 00 ;ynl288w::kanMX4 | OpenBiosystems |
| 7 | ccr4 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 30 ; yal021c::kanMX4 | Generated in the lab |
| 8 | cth1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; $y d r 151 c::$ kanMX4 | OpenBiosystems |
| 9 | des1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; vlr270w: :kanMX4 | OpenBiosystems |
| 10 | dcs2 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met 15 40 ; ura3 30 ; yor173w::kanMX4 | OpenBiosystems |
| 11 | dhh1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 10 ; ura3 30 ; ydl160c::kanMX4 | Generated in the lab |
| 12 | dom34 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; ynl001w::kanMX4 | OpenBiosystems |
| 13 | edc 1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; ygl222c::kanMX4 | OpenBiosystems |
| 14 | edc2 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; yer035w: :kanMX4 | Generated in the lab |
| 15 | edc3 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; yel015w:: kanMX4 | OpenBiosystems |
| 16 | esc1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; <br> met1500; ura3 30 ; ymr219w: kanMX4 | OpenBiosystems |
| 17 | hbs1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 0 ; ura3 40 ; ykr084c::kanMX4 | OpenBiosystems |
| 18 | 1sm1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met $15 \Delta 0$; ura $3 \Delta 0$; yjl124c:: kanMX4 | OpenBiosystems |
| 19 | 1sm6 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; $y d r 378 c::$ kanMX4 | OpenBiosystems |
| 20 | 1sm7 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; ynl147w::kanMX4 | OpenBiosystems |
| 21 | not3 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met $15 \Delta 0$; ura $3 \Delta 0$; yil038c::kanMX4 | Generated in the lab |
| 22 | pan2 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 10 ; ura3 00 ; vgl094c::kanMX4 | OpenBiosystems |
| 23 | pan3 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 10 ; ura3 $\Delta 0$; ykl025c::kanMX4 | OpenBiosystems |


| 24 | pat1 | BY4741 MATa his $3 \Delta 1$; leu2 $\Delta 0$; met15 $\Delta 0$; ura3 $\Delta 0$; ycr077c::kanMX4 | Generated in the lab |
| :---: | :---: | :---: | :---: |
| 25 | pop2 | BY4741 MATa his $3 \Delta 1$; leu2 $\Delta 0$; met1500; ura3 30 ; ynr052c::kanMX4 | Generated in the lab |
| 26 | pub1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 40 ; ynl016w::kanMX4 | OpenBiosystems |
| 27 | puf1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; yjr091c::kanMX4 | OpenBiosystems |
| 28 | puf2 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 00 ; ura3 30 ; ypr042c::kanMX4 | OpenBiosystems |
| 29 | puf3 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 0 ; ura3 $\Delta 0 ;$ yll013c:: kanMX4 | OpenBiosystems |
| 30 | puf4 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 $\Delta 0$; ura3 30 ; ygl014w::kanMX4 | OpenBiosystems |
| 31 | puf5 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 0 ; ura3 30 ; ygl178w::kanMX4 | OpenBiosystems |
| 32 | puf6 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 $\Delta 0$; ura3 $\Delta 0$; ydr496c::kanMX4 | OpenBiosystems |
| 33 | rail | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 $\Delta 0$; ura3 $\Delta 0$; ygl246c::kanMX4 | OpenBiosystems |
| 34 | rrp47 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met1540; ura3 30 ; yhr081w::kanMX4 | Generated in the lab |
| 35 | rrp6 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; <br> met1540; ura3 30 ; yor001w::kanMX4 | Generated in the lab |
| 36 | rtt103 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 10 ; ura3 $30 ; y d r 289 c:: k a n M X 4$ | OpenBiosystems |
| 37 | scd6 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met1500; ura3 40 ; ypr129w::kanMX4 | OpenBiosystems |
| 38 | ski2 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 0 ; ura3 $\Delta 0$; ylr398c::kanMX4 | Generated in the lab |
| 39 | ski3 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met1500; ura3 30 ; ypr189w::kanMX4 | OpenBiosystems |
| 40 | ski7 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 50 ; ura3 30 ; yor076c::kanMX4 | OpenBiosystems |
| 41 | ski8 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 $\Delta 0$; ura3 $\Delta 0 ; y g l 213 c:$ :kanMX4 | OpenBiosystems |
| 42 | swt1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 $\Delta 0$; ura3 00 ; yor $166 c:: k a n M X 4$ | OpenBiosystems |
| 43 | tex1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 30 ; ynl253w::kanMX4 | OpenBiosystems |
| 44 | thp2 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; <br> met15 0 ; ura3 $\Delta 0$; yhr167w::kanMX4 | OpenBiosystems |
| 45 | tpal | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 00 ; ura3 30 ; yer049w::kanMX4 | OpenBiosystems |
| 46 | upf2 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met1500; ura3 00 ; yhr077c::kanMX4 | OpenBiosystems |
| 47 | upf3 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 0 ; ura3 $\Delta 0$; ygr072w::kanMX4 | OpenBiosystems |
| 48 | xrn1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; | Generated in the lab |


|  |  | met15 0 ; ura3 0 ; ygl173c::kanMX4 |  |
| :---: | :---: | :---: | :---: |
| 49 | Y40343 | W303 MATalpha tor 1-1 fpr $1:$ :NAT RPL13A-2×FKBP12::TRP1 | Euroscarf |
| 50 | XRN1AA | W303 MATalpha torl-1 fpr $1:$ :NAT RPL13A-2×FKBP 12::TRP1 YGL173C::YGL173C-FRB-GFP-KanMX4 | Generated in the lab |
| 51 | xrn1pm | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 10 ; ura3 $\Delta 0$; ygl173c:: pFA6a-ygl173c-D206A,D208A-3HA-His3MX | Generated in the lab |
| 52 | XRN1-WT | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 10 ; ura3 40 ; YGL173C:: pFA6a-YGL173C-3HA-His3MX | Generated in the lab |
| 53 | nrg1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 40 ; ura3 $\Delta 0$; $y d r 043 c::$ kanMX4 | Generated in the lab |
| 54 | nrg1xrn1 | BY4741 MATa his $3 \Delta 1$; leu $2 \Delta 0$; met15 10 ; ura3 $\Delta 0$; ydr043c::kanMX4; $y g l 173 c:$ :natMX | Generated in the lab |

