Supplementary information for:

Dom34-Hbs1 mediated dissociation of inactive 80S ribosomes promotes restart of translation after stress.

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Running title: Dom34-Hbs1 modulates ribosomal subunit availability after stress

Supplementary Material and Methods

In vitro translation

Translational extracts (Tuite & Plesset, 1986) were prepared from a $dom34\Delta hbs1\Delta$ strain (BSY2550) as follows. 4 L of culture at OD₆₀₀ = 1 was washed in 200 ml cold water, incubated in 100 ml β-mercaptoethanol 10mM; EDTA 2 mM for 30 minutes at room temperature, washed in 100 ml cold sorbitol 1M and resuspended in 1M sorbitol at room temperature at a concentration of 10 ml/g cells, spinning at 2000 x g for 5 min at 4°C in between. Zymolyase was added at 4 μg/ml final concentration and the reaction was stopped by pelleting the resulting spheroplast at 1000 x g for 10 min at room temperature when 75% of cells had converted to spheroplasts. Spheroplasts were washed in 200 ml sorbitol 1.2 M and incubated in 500 ml YPDA-sorbitol 1M at 25°C 40 rpm. Spheroplasts were harvested at 1000 x g for 10 min at 4°C and lysed using glass beads (0.5 ml/g cells) in lysis buffer (20 mM Hepes-KOH pH7.4; 100 mM KOAc; 2 mM Mg(OAC); 2 mM DTT; 0.5 mM PMSF; protease

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inhibitor cocktail) (1 ml/g cells) in 5 cycles of shaking vigorously at 2 Hz for 20 seconds with 1 minute intervals on ice. Lysates were cleared spinning 30 000 x g for 15 min at 4°C, then 100 000 x g for 30 min at 4°C. Glycerol was added at 10% final concentration for storage at -80°C.

In vitro translation was performed basically as described in (Tarun & Sachs, 1995). Translational extracts were incubated with 150 U/ml micrococcal nuclease (New England Biolabs) in presence of 480 µM CaCl₂ for 5 minutes at 26°C, before adding 2 mM EGTA on ice. 7.5 µl extract was added to 7.5 µl translation mix containing 0,1 µl RNasin (Promega), 500 ng firefly luciferase-A(50) mRNA (Gallie et al, 1991) and 4 µg creatine phosphokinase (Roche) in 22 mM Hepes-KOH pH 7.4; 120 mM KOAc; 2 mM MgOAc; 750 µM ATP; 100 μM GTP; 25 mM creatine phosphate; 40 μM amino acid mixture (Promega); 1.7 mM DTT. After 1 h incubation at 26°C luciferase activity was measured in 10 s measurements using a Lumat LB 9507 luminometer (Berthold technologies) adding 1 µl translation reaction to 50 µl luciferine mix (470 µM luciferine; 530 µM ATP; 270 µM coenzyme A; 20 mM Trisphosphate pH 7.8; 1.07 mM MgCl₂; 2.7 mM MgSO₄; 100 μM EDTA; 33.3 mM DTT). Dom34 added to the translation reaction was purified as described in (Shoemaker et al., 2010; Shoemaker & Green, 2011). Hbs1 was purified from BL21 codon+ cells expressing Cterminally 6xHis-tagged Hbs1 grown in Autoinduction media Terrific Broth Base including Trace elements (Formedium) over Ni-NTA agarose (Qiagen) after cell lysis in a Cell Disruptor (Constant Systems) at 1.55 kbar, using lysis buffer (75 mM Hepes pH 7.9, 300 mM NaCl, 5 mM β-mercaptoethanol, 1% Tween, 20 mM imidazole, 10% glycerol, 2 mM MgCl₂), wash buffer (50 mM Hepes pH 7.9, 500 mM NaCl, 5 mM β-mercaptoethanol, 20 mM imidazole, 10% glycerol, 2 mM MgCl₂) and elution buffer (50 mM Hepes pH 7.9, 500 mM NaCl, 5 mM β-mercaptoethanol, 300 mM imidazole, 10% glycerol, 2 mM MgCl₂). Both Hbs1 and the Dom34-Hbs1 complex were further purified over a Superdex 75 10/300 GL column (GE Healthcare) in 20mM Tris-Cl pH 7.5, 200mM NaCl, 5mM β-mercaptoethanol, 5% glycerol.

Yeast growth curves

Yeast was grown to OD_{600} 0.6 in YPDA, shifted to the indicated temperature and grown for another 2 hours. The rest of the experiment was done at the indicated temperatures. Cultures were split into 2 x 10 ml and resuspended in 10 ml YPA or YPDA for 10 minutes. Then 3 ml of each cultures was pelleted and resuspended in 15 ml YPDA (to an OD_{600} of \sim 0.2). OD_{600}

was measured at the indicated time points. OD_{600} series corresponding to the same cultures were normalized for the OD_{600} at time = 0 minutes to be 0.2.

Supplementary table 1: Strains and plasmids

Description	Reference
(genotype for strains; vector/insert/marker for plasmids)	
$MAT \alpha$, $ura31$, $trp1\Delta$, $ade21$, $leu23$, $leu33$, l	(Baudin-Baillieu
	et al, 1997)
$MAT \alpha$, $ura31$, $trp1\Delta$, $ade21$, $leu23$, $l12$, $his311$, $l5$,	(van den Elzen
dom34∆::HIS3	et al, 2010)
<i>MAT a, ura31, trp1∆, ade 21, leu23,112, his311,15,</i>	(van den Elzen
hbs1\Delta::KanR	et al, 2010)
MAT a, ade 2-1 his3-11,15, leu2-3,112, trp1delta ura3-1,	This study
$dom34\Delta$::HIS3, $hbs1\Delta$::KanR	
$MAT \ a, \ ura3-1, \ trp1\Delta, \ ade2-1, \ leu2-3,112, \ his3-11,15,$	Gift from F.
$stm1\Delta$::TRP1	Lacroute and F. Wyers
MAT a, ura3-1, trp1 Δ , ade2-1, leu2-3,112, his3-11,15, stm1 Δ ::TRP1, dom34 Δ ::HIS3	This study
"DC415/ HDC1 DDOTEIN 4 magazines LEU2	(van den Elzen
pRS413/ HBS1-PROTEIN A + promoter/LEU2	et al, 2010)
pBS3675 pRS415/ hbs1 V176G-PROTEIN A + promoter/LEU2	(van den Elzen
	et al, 2010)
pBS3685 pRS415/ DOM34-3HA + promoter/LEU2	(van den Elzen
	et al, 2010)
pBS3701 pRS415/ dom34 E361R-3HA + promoter/LEU2	(van den Elzen
	et al, 2010)
pRS415/ hbs1ΔN-ter (2-149)-PROTEIN A + promoter/LEU2	This study
	(genotype for strains; vector/insert/marker for plasmids) MAT α, ura31, trp1Δ, ade21, leu23,112, his311,15 MAT α, ura31, trp1Δ, ade21, leu23,112, his311,15, dom34Δ::HIS3 MAT a, ura31, trp1Δ, ade 21, leu23,112, his311,15, hbs1Δ::KanR MAT a, ade 2-1 his3-11,15, leu2-3,112, trp1delta ura3-1, dom34Δ::HIS3, hbs1Δ::KanR MAT a, ura3-1, trp1Δ, ade2-1, leu2-3,112, his3-11,15, stm1Δ::TRP1 MAT a, ura3-1, trp1Δ, ade2-1, leu2-3,112, his3-11,15, stm1Δ::TRP1, dom34Δ::HIS3 pRS415/ HBS1-PROTEIN A + promoter/LEU2 pRS415/ bbs1 V176G-PROTEIN A + promoter/LEU2 pRS415/ DOM34-3HA + promoter/LEU2 pRS415/ dom34 E361R-3HA + promoter/LEU2