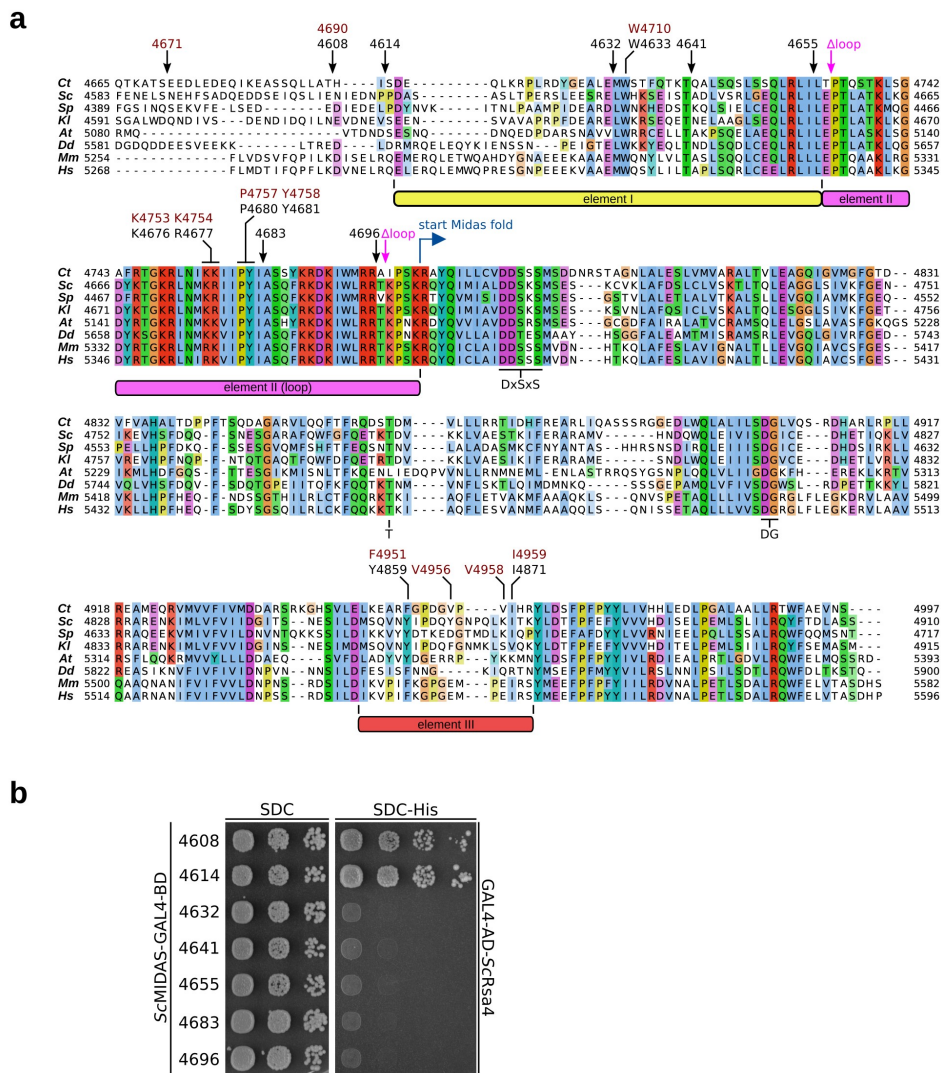


Crystal structures of Rea1-MIDAS bound to its ribosome assembly factor ligands resembling integrin–ligand-type complexes

Yasar Luqman Ahmed et al.

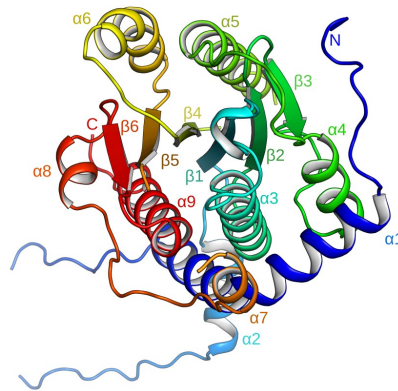
Supplementary Information

Supplementary Figures

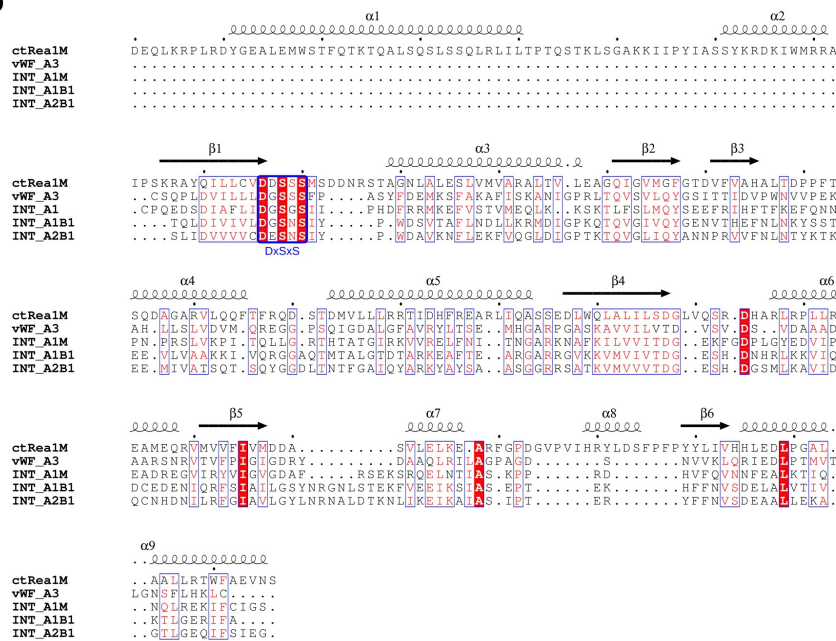


Supplementary Figure 1 Multiple sequence alignment of the Rea1-MIDAS domain and truncation analysis. (a) Multiple sequence alignment of the Rea1 with homologs showing the MIDAS domain. The sequences of *Chaetomium thermophilum* (Ct), *Saccharomyces cerevisiae* (Sc), *Schizosaccharomyces pombe* (Sp), *Kluyveromyces lactis* (Kl), *Arabidopsis thaliana* (At), *Dictyostelium discoideum* (Dd), *Mus musculus* (Mm), and *Homo sapiens* (Hs) were aligned with Clustal Omega and visualized with Jalview. The Rea1-specific elements (I–III) and the residues required for coordination of the Mg²⁺ ion are highlighted below the alignment. Indicated above the alignment are the generated truncations (arrows) and point mutations in *C.thermophilum* (red) and *S.cerevisiae* (black). The boundaries of the Δloop constructs are indicated in pink. (b) Yeast two-hybrid analysis of the interactions between the indicated MIDAS constructs with full-length Rsa4 from *S. cerevisiae*. The Rsa4 construct was fused to an N-terminal GAL4-AD (activation domain) and the MIDAS constructs were fused to a C-terminal GAL4-BD (binding domain). The residue numbers indicate the N-terminal boundary of the MIDAS constructs. Plasmids were co-transformed into yeast (PJ69-4A) and transformants were spotted in tenfold serial dilutions on SDC (SDC–Leu–Trp) and SDC–His (SDC–Leu–Trp–His) selective plates. Cell growth was monitored after incubation for 3 days at 30 °C.

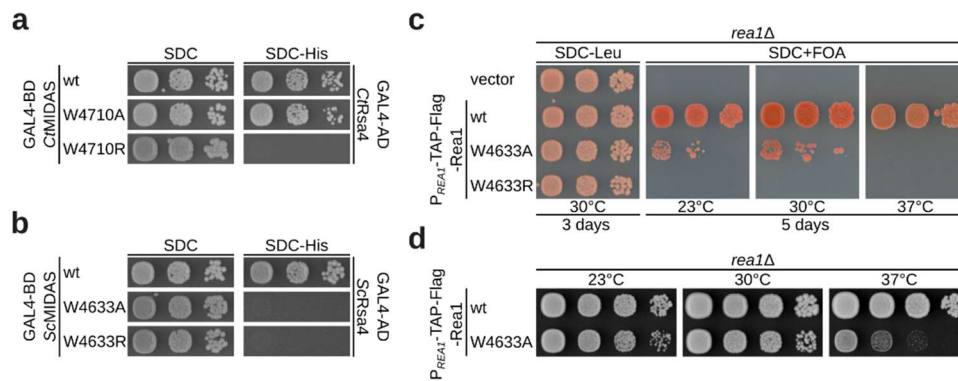
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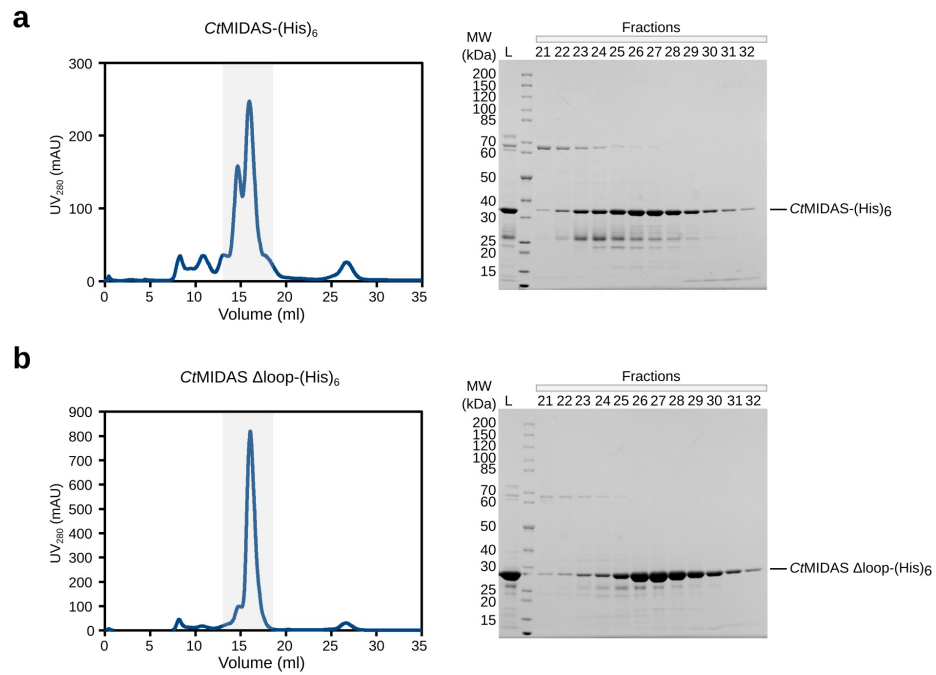
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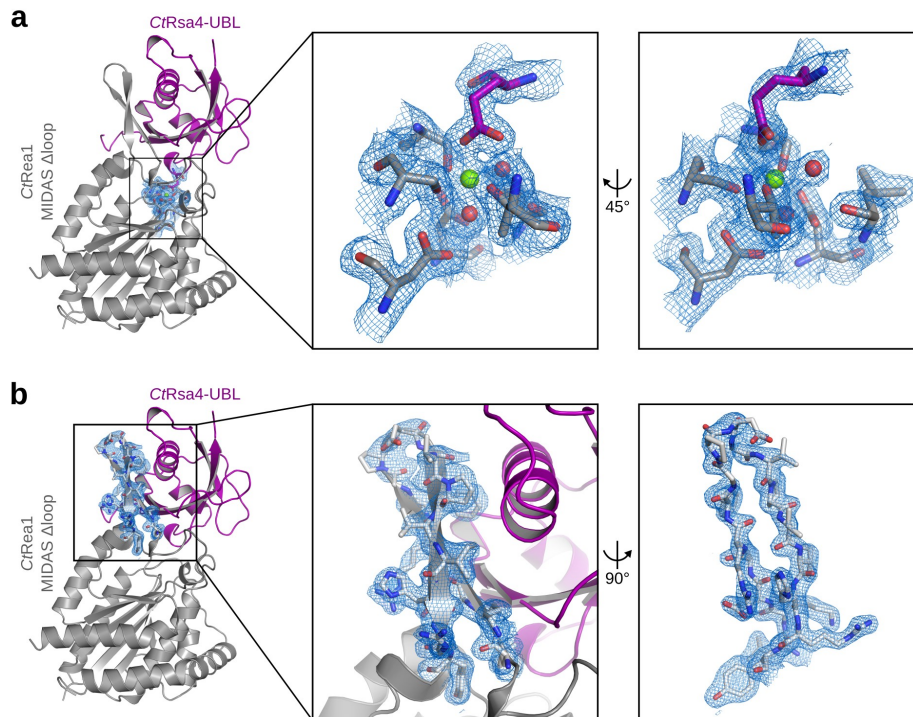
Supplementary Figure 2. Secondary structure comparison of the Rea1-MIDAS and integrin structures. (a) Structure of the Rea1-MIDAS domain from *C. thermophilum*. (b) Sequence and structure alignment between *CtRea1* and depicted integrin structures. The alignment was created with MAFFTASH. The following structures were aligned: *ctRea1M* (*C. thermophilum* Rea1-MIDAS, sequence contains gaps not covered in the structure), *vWF_A3* (von Willebrand factor A3 domain, PDB ID: 1AO3), *INT_A1M* (integrin alpha M I domain, PDB ID: 1MF7), *INT_A1B1* (integrin alpha1beta1 I-domain, PDB ID: 1QCY) and *INT_A2B1* (integrin alpha2beta1 I-domain, PDB ID: 5HJ2).



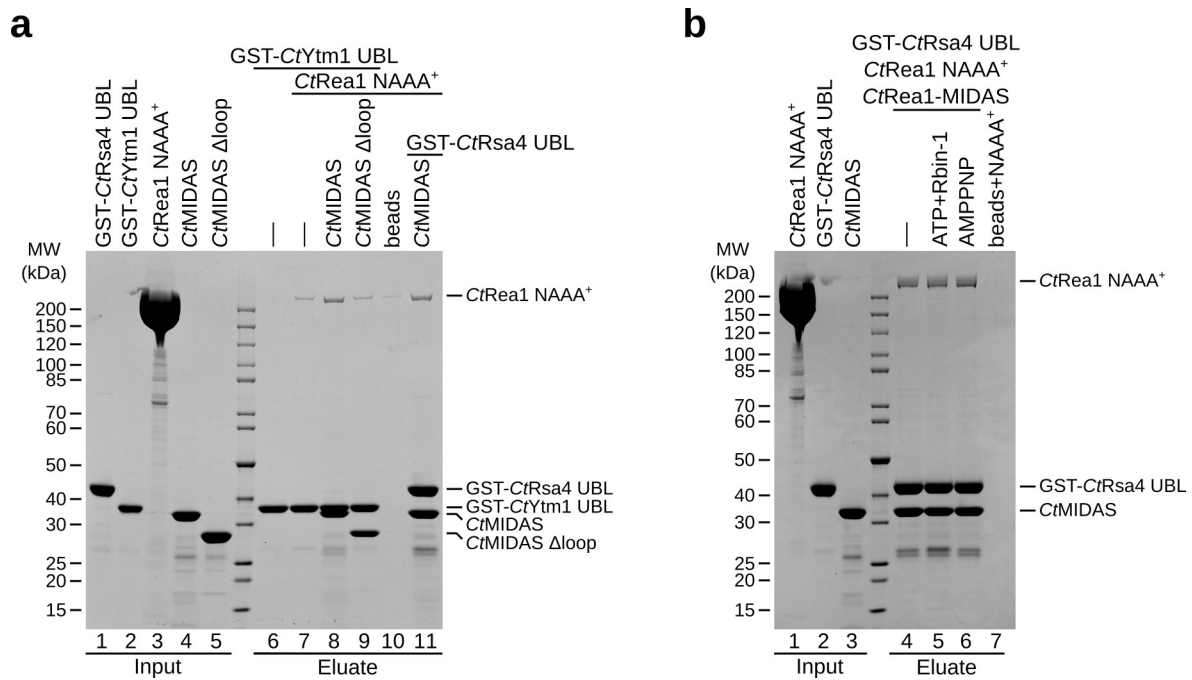
Supplementary Figure 3. Functional analysis of the conserved Try residue in the N-terminal extension of Rea1-MIDAS. (a,b) Yeast two-hybrid analysis of the Rea1-MIDAS and Rsa4 constructs from *C. thermophilum* (a) and *S. cerevisiae* (b). The MIDAS constructs were fused to an N-terminal GAL4-BD and the Rsa4 constructs to an N-terminal GAL4-AD. Transformants were spotted on SDC (SDC–Leu–Trp) plates or SDC–His (SDC–Leu–Trp–His) selective plates and cell growth was monitored after incubation for 3 days at 30 °C. (c) An empty plasmid control or the indicated Rea1 constructs, fused to an N-terminal TAP-Flag tag, were transformed into a *rea1Δ* shuffle strain. Cells were spotted in 10-fold serial dilutions on SDC–Leu and SDC+FOA plates and cells were grown at the indicated temperatures for 3 and 5 days, respectively. (d) Growth analysis of the indicated Rea1 constructs after shuffling on SDC+FOA plates. Cells were spotted in tenfold serial dilutions on YPD plates and growth at the indicated temperatures was monitored after 2 days.



Supplementary Figure 4. Purification of CtMIDAS and CtMIDAS Δ loop. (a,b) The left-hand panels show the size-exclusion chromatography profiles of CtMIDAS-(His)₆ (amino acids 4690-4997) (a) and CtMIDAS Δ loop-(His)₆ (amino acids 4690-4733-GSG-4774-4997) (b). The absorbance at 280 nm (mAU) was plotted against the elution volume (ml). Fractions highlighted in gray were analyzed by SDS-PAGE and Coomassie staining (right-hand panels).



Supplementary Figure 5. Structure of the CtRea1-MIDAS Δ loop-CtRsa4-UBL complex. (a,b) The CtRea1-MIDAS Δ loop-CtRsa4-UBL model and corresponding electron densities of the Mg²⁺ coordination site (a) and the β -hairpin of the Rea1-specific element III (b) contoured at 1.5 and 1.0 sigma, respectively. The CtRsa4-UBL is highlighted in purple and the CtRea1-MIDAS Δ loop is shown in gray. Close-up views are shown in two orientations.



Supplementary Figure 6. The MIDAS element II loop is required for interaction of the Rea1-MIDAS–Ytm1-UBL complex with the Rea1-AAA⁺ ring. (a) GST-tagged CtYtm1-UBL or CtRsa4-UBL were incubated with CtRea1-MIDAS or CtRea1-MIDAS Δ loop and the CtRea1-NAAA⁺ ring domain. After washing, the GST-tagged CtYtm1- and CtRsa4-UBL domains and bound material were eluted with GSH and analyzed with SDS-PAGE and Coomassie staining. Protein inputs are shown in lanes 1–5, final eluates in lanes 6–11. **(b)** Binding assay between GST-tagged CtRsa4-UBL, CtRea1-MIDAS and Ct-NAAA⁺ ring domain in the absence of nucleotides, in presence of ATP (2 mM final concentration) and the inhibitor Rbin-1 (0.1 mM final concentration), or in presence of non-hydrolysable AMPPNP (2 mM final concentration). GST-CtRsa4-UBL and bound material was eluted with GSH and analyzed with SDS-PAGE and Coomassie staining. Protein inputs are shown in lanes 1–3, final eluates in lanes 4–7.

Supplementary Table 1. *E. coli* expression plasmids used in this study

Name	Relevant information	Source
pET24d-CfMIDAS-(His) ₆	Kan ^R , T7 promoter, <i>lac</i> operator, (aa 4671–4997)	This study
pETHis-(His) ₆ -CfYtm1-UBL	Kan ^R , T7 promoter, <i>lac</i> operator, (aa 8–98)	This study
pETHis-(His) ₆ -CfRsa4-UBL	Kan ^R , T7 promoter, <i>lac</i> operator, (aa 31–128)	This study
pET24d-(His) ₆ -GST-CfRsa4-UBL	Kan ^R , T7 promoter, <i>lac</i> operator, (aa 1–128)	This study
pET24d-(His) ₆ -GST-CfYtm1-UBL	Kan ^R , T7 promoter, <i>lac</i> operator, (aa 10–98)	This study
pET-15b-CfMIDAS-(His) ₆	Amp ^r , T7 promoter, <i>lac</i> operator, (aa 4690–4997)	This study
pET-15b-CfMIDASΔloop-(His) ₆	Amp ^r , T7 promoter, <i>lac</i> operator, (aa 4690–4733-GSG-4774–4997)	This study
pET24d-GST-CfKap104	Kan ^R , T7 promoter, <i>lac</i> operator	1
pET-15b-CfSyo1-(His) ₆	Amp ^r , T7 promoter, <i>lac</i> operator	1
pET-15b-CfMIDAS PY>A-(His) ₆	Amp ^r , T7 promoter, <i>lac</i> operator, (aa 4690–4997)	This study
pET-15b-CfMIDAS KK>A-(His) ₆	Amp ^r , T7 promoter, <i>lac</i> operator, (aa 4690–4997)	This study
pET-15b-CfMIDAS PY/KK>A-(His) ₆	Amp ^r , T7 promoter, <i>lac</i> operator, (aa 4690–4997)	This study

Supplementary Table 2. *S. cerevisiae* plasmids used in this study

Name	Relevant information	Source
pVA3-1 (p53)	2 μ , <i>TRP1</i> , GAL4 BD-murine p53 (aa 72–390), pGBT9	Takara Bio Inc.
pTD1-1 (SV40)	2 μ , <i>LEU2</i> , GAL4 AD-SV40 large T-antigen (aa 84–708), pACT2	Takara Bio Inc.
YCplac111	CEN, <i>LEU2</i>	2
pGADT7-ScRSA4	2 μ , <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-AD	3
pG4BDC22-Gly-aa4608-ScMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal GAL4-BD	This study
pG4BDC22-aa4614-ScMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal GAL4-BD	This study
pG4BDC22-Gly-aa4632-ScMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal GAL4-BD	This study
pG4BDC22-aa4641-ScMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal GAL4-BD	This study
pG4BDC22-Gly-aa4655-ScMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal GAL4-BD	This study
pG4BDC22-aa4683-ScMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal GAL4-BD	This study
pG4BDC22-aa4696-ScMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal GAL4-BD	This study
pGADT7-CfRSA4	2 μ , <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-AD	4
pG4BDN22-aa4690-CfMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS Δ loop	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS DAA	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS Δ loop DAA	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4608-ScMIDAS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4608-ScMIDAS Δ loop	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4608-ScMIDAS DAA	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4608-ScMIDAS Δ loop DAA	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS F4951R	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS F4951A	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS I4959R	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS I4959A	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS V4956R	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS V4956A	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS V4958R	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4690-CfMIDAS V4958A	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDC22-Gly-aa4608-ScMIDAS Y4859R	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDC22-Gly-aa4608-ScMIDAS Y4859A	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDC22-Gly-aa4608-ScMIDAS I4871R	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDC22-Gly-aa4608-ScMIDAS I4871A	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDC22-Gly-aa4608-ScMIDAS Y4859R I4871R	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDC22-Gly-aa4608-ScMIDAS Y4859A I4871A	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
YCplac111-PREA1-TAP-Flag-REA1	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	5
YCplac111-PREA1-TAP-Flag- <i>rea1</i> Y4859A	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> Y4859R	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> I4871A	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> I4871R	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> Y4859A I4871A	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> Y4859R I4871R	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
pRS314-RSA4	CEN, <i>TRP1</i> , <i>PRSA4</i> , <i>TRSA4</i> ,	3
pRS314- <i>rsa4</i> -1	CEN, <i>TRP1</i> , <i>PRSA4</i> , <i>TRSA4</i> ,	3
pRS314-YTM1	CEN, <i>TRP1</i> , <i>PYTM1</i> , <i>TYTM1</i> ,	6
pRS314- <i>ytm1</i> S78L	CEN, <i>TRP1</i> , <i>PYTM1</i> , <i>TYTM1</i> ,	6
YCplac111-PREA1-TAP-Flag- <i>rea1</i> Δ MIDAS	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag (aa 1–4621)	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> Δ loop	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study

Name	Relevant information	Source
YCplac111-PREA1-TAP-Flag- <i>rea1</i> DAA	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	5
YCplac111-PGAL1-10-TAP-Flag- <i>REA1</i>	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	5
YCplac111-PGAL1-10-TAP-Flag- <i>rea1</i> ΔMIDAS	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PGAL1-10-TAP-Flag- <i>rea1</i> Δloop	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PGAL1-10-TAP-Flag- <i>rea1</i> DAA	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PREA1-GFP- <i>REA1</i>	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal GFP tag	5
YCplac111-PREA1-GFP- <i>rea1</i> DAA	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal GFP tag	5
YCplac111-PREA1-GFP- <i>rea1</i> ΔMIDAS	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal GFP tag	This study
YCplac111-PREA1-GFP- <i>rea1</i> Δloop	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal GFP tag	This study
YCplac111-PADH1-ScMIDAS loop-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	This study
YCplac111-PADH1-CtMIDAS loop-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	This study
YCplac111-PADH1-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	1
YCplac111-PADH1-CtMIDAS loop PY>A-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	This study
YCplac111-PADH1-CtMIDAS loop KK>A-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	This study
YCplac111-PADH1-CtMIDAS loop PY/KK>A-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	This study
YCplac111-PADH1-ScMIDAS loop PY>A-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	This study
YCplac111-PADH1-ScMIDAS loop KR>A-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	This study
YCplac111-PADH1-ScMIDAS loop PY/KR>A-3xGFP	CEN, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , C-terminal 3xGFP tag	This study
pG4BDN22-aa4608-ScMIDAS loop>L4-PY-NLS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4608-ScMIDAS loop>Hrp1-PY-NLS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
pG4BDN22-aa4608-ScMIDAS loop>Syo1-PY-NLS	CEN, <i>TRP1</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal GAL4-BD	This study
YCplac111-PREA1-GFP- <i>rea1</i> loop>L4-PY-NLS	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal GFP tag	This study
YCplac111-PREA1-GFP- <i>rea1</i> loop>Syo1-PY-NLS	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal GFP tag	This study
YCplac111-PREA1-GFP- <i>rea1</i> loop>Hrp1-PY-NLS	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal GFP tag	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> loop>L4-PY-NLS	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> loop>Syo1-PY-NLS	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PREA1-TAP-Flag- <i>rea1</i> loop>Hrp1-PY-NLS	CEN, <i>LEU2</i> , <i>PREA1</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PGAL1-10-TAP-Flag- <i>rea1</i> loop>L4-PY-NLS	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PGAL1-10-TAP-Flag- <i>rea1</i> loop>Syo1-PY-NLS	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PGAL1-10-TAP-Flag- <i>rea1</i> loop>Hrp1-PY-NLS	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal TAP-Flag tag	This study
YCplac111-PGAL1-10-GFP- <i>REA1</i>	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal GFP tag	5
YCplac111-PGAL1-10-GFP- <i>rea1</i> DAA	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal GFP tag	5
YCplac111-PGAL1-10-GFP- <i>rea1</i> loop>Hrp1-PY-NLS	CEN, <i>LEU2</i> , <i>PGAL1-10</i> , <i>TREA1</i> , N-terminal GFP tag	This study
pADH181-ProtA-TEV-NAAA ⁺ CtRea1	2μ, <i>LEU2</i> , <i>PADH1</i> , <i>TADH1</i> , N-terminal ProtA-TEV tag (aa 1–2390)	This study

Supplementary Table 3. *S. cerevisiae* strains used in this study

Name	Relevant genotype	Source
PJ69-4A	<i>trp1-901, leu2-3,112, ura3-52, his3-200, gal4Δ, gal80Δ, LYS2::GAL1-⁷HIS3, GAL2-ADE2, met2::GAL7-lacZ</i>	
W303	Wild type, MAT α	8
DS1-2b *	Wild type	9
<i>rea1Δ</i> shuffle	<i>rea1::kanMX6, pRS416-REA1</i>	3
<i>rea1Δ rsa4Δ</i> shuffle	<i>rea1::kanMX6, rsa4::His3MX6, pRS416-REA1, pRS316-RSA4</i>	3
<i>rea1Δ ytm1Δ</i> shuffle	<i>rea1::His3MX6, ytm1::hphNT1, pRS416-REA1, pRS316-YTM1</i>	6
Aid-HA-REA1	<i>P_{REA1}-Aid-HA::natNT2, P_{ADH1}-OsTIR1-9xmyc::TRP1</i>	This study
RIX1-TAP RPL3-Flag *	<i>RIX1-TAP::TRP1, RPL3-Flag::natNT2</i>	5

Supplementary References

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