

**Fig. S1. Toxicity of A** $\beta$  **alone.** (A) Ten-fold dilutions of exponentially growing cultures of BY4742 cells transformed with plasmids carrying the different chimeric constructions under the GAL10 promoter were spotted onto SD (-) or SG (+) agar supplemented with 20 mg/l histidine, 20 mg/l lysine and 60 mg/l leucine. The cells were incubated at 30°C for 3 days. (B) BY4742 cells expressing the different chimeric proteins (6 hours of expression) were collected for total-protein extracts. Equal quantities of proteins were separated by SDS-PAGE on a 12% polyacrylamide gel, transferred onto a nitrocellulose membrane and exposed to anti-A $\beta$  (Tebu) antibodies.

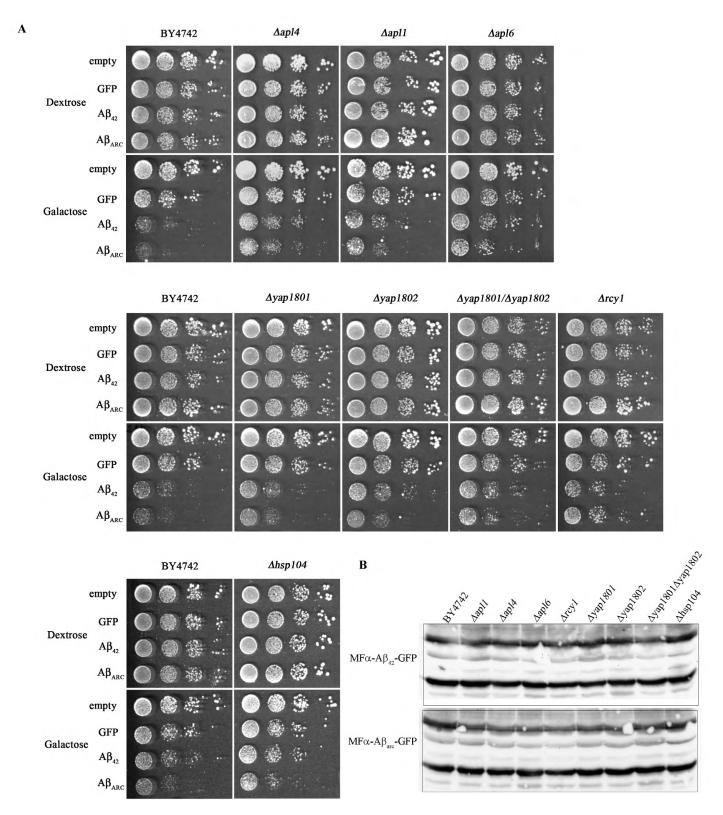


Fig. S2. Yeast mutants modulate A $\beta$  toxixity without changing its protein level. (A) Ten-fold dilutions of exponentially growing cultures of deleted strains transformed with plasmids carrying the different chimeric constructions under the GAL10 promoter were spotted on the same plate onto SD (-) or SG (+) agar supplemented with 20 mg/l histidine, 20 mg/l lysine and 60 mg/l leucine. (B) Cells expressing the different chimeric proteins (6 hours of expression) were collected for total-protein extracts. Equal quantities of proteins were separated by SDS-PAGE on a 12% polyacrylamide gel, transferred onto a nitrocellulose membrane, and then exposed to monoclonal anti-GFP antibodies (Sigma).

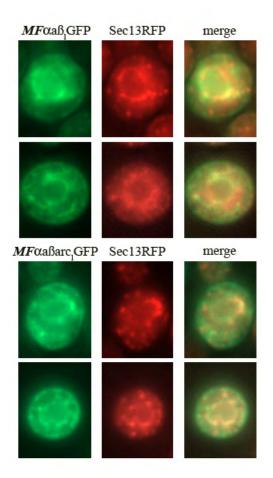


Fig. S3. A $\beta$ -GFP species colocalize with ER. Wild-type or RFP-tagged strains were grown for 6 hours in SG liquid medium supplemented with 0.67% casaminoacids to induce the expression of the chimeric proteins and were then examined by epifluorescence microscopy.

Table 1. Yeast strains used in this study

| Name of strain    | Genotype  | Source                          |
|-------------------|---|---------------------------------|
| BY4742            | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0$                                      | Eurocarf yeast deletion library |
| $\Delta a p l 4$  | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0, YPR029C::KanMX4$                     | Eurocarf yeast deletion library |
| $\Delta apl1$     | $MATa, his3\Delta 1, leu2\Delta 0, ura3\Delta 0, YJR005W::KanMX4$                     | Eurocarf yeast deletion library |
| $\Delta apl2$     | $MATa$ , his $3\Delta 1$ , leu $2\Delta 0$ , ura $3\Delta 0$ , YKL135C::KanMX4        | Eurocarf yeast deletion library |
| Δapl3             | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0, YBL037W ::KanMX4$                    | Eurocarf yeast deletion library |
| $\Delta apm1$     | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0, YPL259C::KanMX4$                     | Eurocarf yeast deletion library |
| $\Delta apm4$     | $MATa, his3\Delta 1, leu2\Delta 0, ura3\Delta 0, YOL062C::KanMX4$                     | Eurocarf yeast deletion library |
| Δgga2             | $MATa$ , his $3\Delta 1$ , leu $2\Delta 0$ , ura $3\Delta 0$ , YHR $108W$ ::Kan $MX4$ | Eurocarf yeast deletion library |
| $\Delta ent5$     | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0, YDR153C::KanMX4$                     | Eurocarf yeast deletion library |
| ∆clc1             | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0, YGR167W::KanMX4$                     | Eurocarf yeast deletion library |
| $\Delta rcyl$     | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0, YJL204C::KanMX4$                     | Eurocarf yeast deletion library |
| Δyap1801          | $MATa$ , his $3\Delta 1$ , leu $2\Delta 0$ , ura $3\Delta 0$ , YHR161C::KanMX4        | Eurocarf yeast deletion library |
| Δyap1802          | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0, YGR241C::KanMX4$                     | Eurocarf yeast deletion library |
| $\Delta hsp104$   | $MATa, his3\Delta 1, leu2 \Delta 0, lys2 \Delta 0, ura3\Delta 0, YLL026W::KanMX4$     | Eurocarf yeast deletion library |
| SEC13-RFP         | $MATa, his3\Delta I, leu2 \Delta 0, lys2 \Delta 0, ura3\Delta 0, YLR208W-RFP-KanMX6$  | Peter Arvidson                  |
| Δyap1801/Δyap1802 | $MATa, his3\Delta I, leu2\Delta 0, ura3\Delta 0, YHR161C::KanMX4, YGR241C::KanMX4$    | This study                      |

## Table S1. Yeast strains used in this study.

## Table S2. Oligonucleotides used in this study

| Number | Sequence   |
|--------|--|
| 792    | AAATACACACACTAAATTACCGGATCCTATGGATGCAGAATTCCGACATG   |
| 794    | ACCAGTGAATAATTCTTCACCTTTAGACATCGCTATGACAACACCGCCCACC |
| 705    | GGATGGCCAGGCAACTTTAG                                 |
| 856    | GAATAATTCTTCACCTTTAGACATAGCTTCAGCCTCTCTTTTATC        |
| 858    | GAATAATTCTTCACCTTTAGACATGGATCCGGTAATTTAGTGTGT        |
| 859    | GTCATGTCGGAATTCTGCATCCATGGATCCGGTAATTTAGTGTGT        |
| 706    | TTTACACTTTATGCTTCCGG                                 |
| 857    | ATGTCTAAAGGTGAAGAATTATTC                             |
| 860    | ATGGATGCAGAATTCCGACATG                               |
|        |  |

Table S2. Oligonucleotides used in this study.