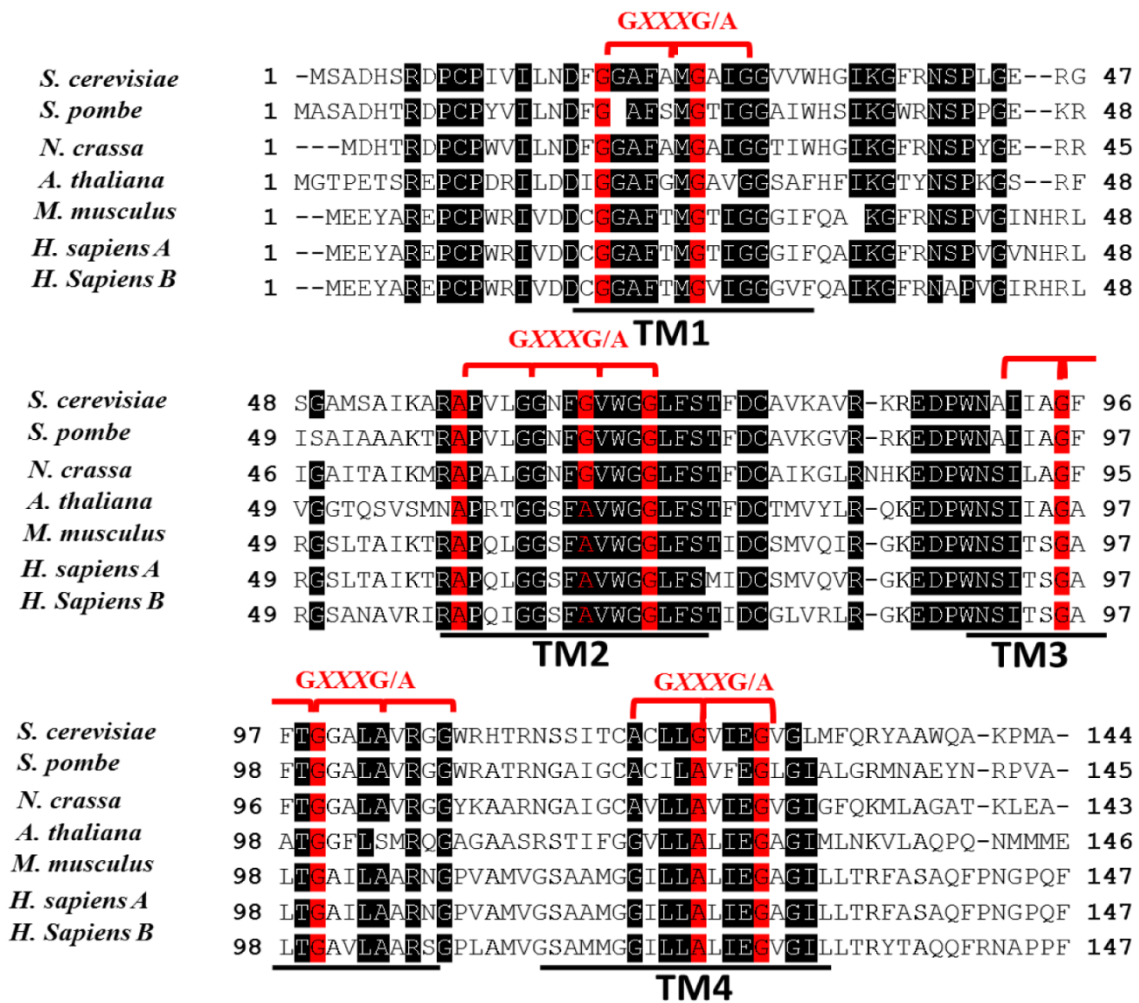


## Supplemental material



**Fig. S1** Sequence alignment of Tim17 protein from different species. (A) Orthologous Tim17 sequences from different species including *S. cerevisiae*, *N. crassa*, *S. pombe*, *A. thaliana*, *M. musculus* and *H. sapiens* were aligned using ClustalW2 online tool. The position of transmembrane helices TM1-TM4 has been indicated. The sequence corresponding to the G/AXXXG/A motif and the position of the conditional mutants isolated are highlighted in red colour.

**Table S1:** Yeast strains used in this study.

Strain	Genotype	Source
TIM17-316	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS316- <i>Tim17</i> ]	(Schilke et al., 2012)
TIM17-314	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> ]	This study
TIM17	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G19A</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G19A</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>A23L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>A23L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G25L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G25L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G19A/A23L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G19A/A23L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G29L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G29L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>A58L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>A58L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G62L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G62L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G66L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G66L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G70L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G70L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>A58L/G70L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>A58L/G70L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>A91L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>A91L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G95L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G95L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G99L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G99L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>A103L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>A103L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G107L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G107L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G123L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G123L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G127L</sub>	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS314- <i>Tim17</i> <sub>G127L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study
Tim17 <sub>G123L/G127L</sub> - TEF	<i>MATa ade2 his3 ura3 leu2 trp1 can1 Tim17Δ::CgHIS3</i> [pRS414- <i>Tim17</i> <sub>G123L/127L</sub> ]- <i>Tim23-FLAG::kanMX4</i>	This study