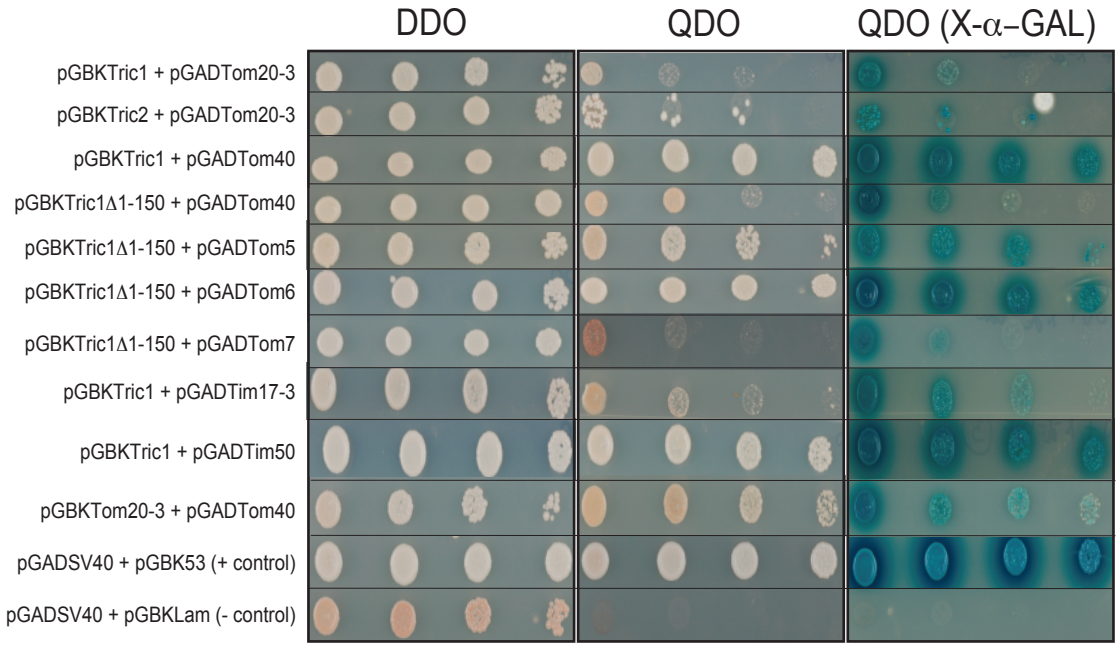
A



B

	Tric1	Tric2	Tric1 Δ1-150	Tom20-3	P53	Lam	pGBKT7 empty
Tim17-1							
Tim17-2							
Tim17-3	X						
Tim23-1							
Tim23-2							
Tim23-3							
Tim21-3							
Tim44							
Tim50	X						
Tom20-2							
Tom20-3	X	X					
Tom40	X		X	X			
Tom5			X				
Tom6			X				
Tom7-1			X				
MPP-a							
RISP							
B14.7							
Cyt C							
SV40					X		
pGADT7 empty							

C

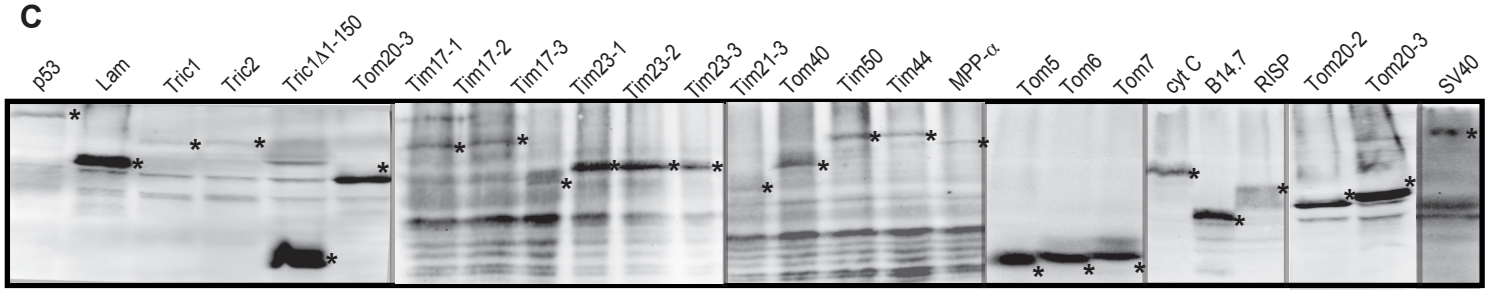


Figure S3. Y2H interactions of Tric with subunits of the protein import apparatus. A. Positive Yeast-2-Hybrid interaction assays of Tric1, Tric2 and the SAM domain of Tric1 (Δ1-150) cloned into pGBK and transformed into AH109 mated agianst various subunits of the TOM and TIM17:23 complex cloned into pGAD and transformed into Y187. Successful diploid mating was determined by growth on Double Drop-Out media (DDO, SD-leu-trp) whilst positive interactors were determined by growth on Quadruple Drop-Out media (QDO, SD-leu-trp-ade-his) with or without the addition of X-α-GAL. **B**. Table representing all the interactions that were tested with pGBKT7 clones listed on the top and pGADT7 clones listed on the right. Positive interactors are indicated by X. **C**. Immunodetection of all yeast transformants to confirm protein expression as detected by anti-c-myc (pGBKT7) and anti-HA (pGADT7). The band corresponding to the specific protein is indicated by a *.