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Author Correction: Loss of mRNA surveillance pathways results in widespread protein aggregation

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The original version of this Article contained errors. Panels upf1 and dom34 in Figure 1 looked to have originated from the same sample. The Authors now reviewed the original data and for clarity all representative images in Figure 1 have been replaced. Additionally, the Authors recalculated the results shown in Figure 1B using the original data and the graph has also been updated. The original Figure 1 is shown below, for reference:

The original version of the Article has been corrected.

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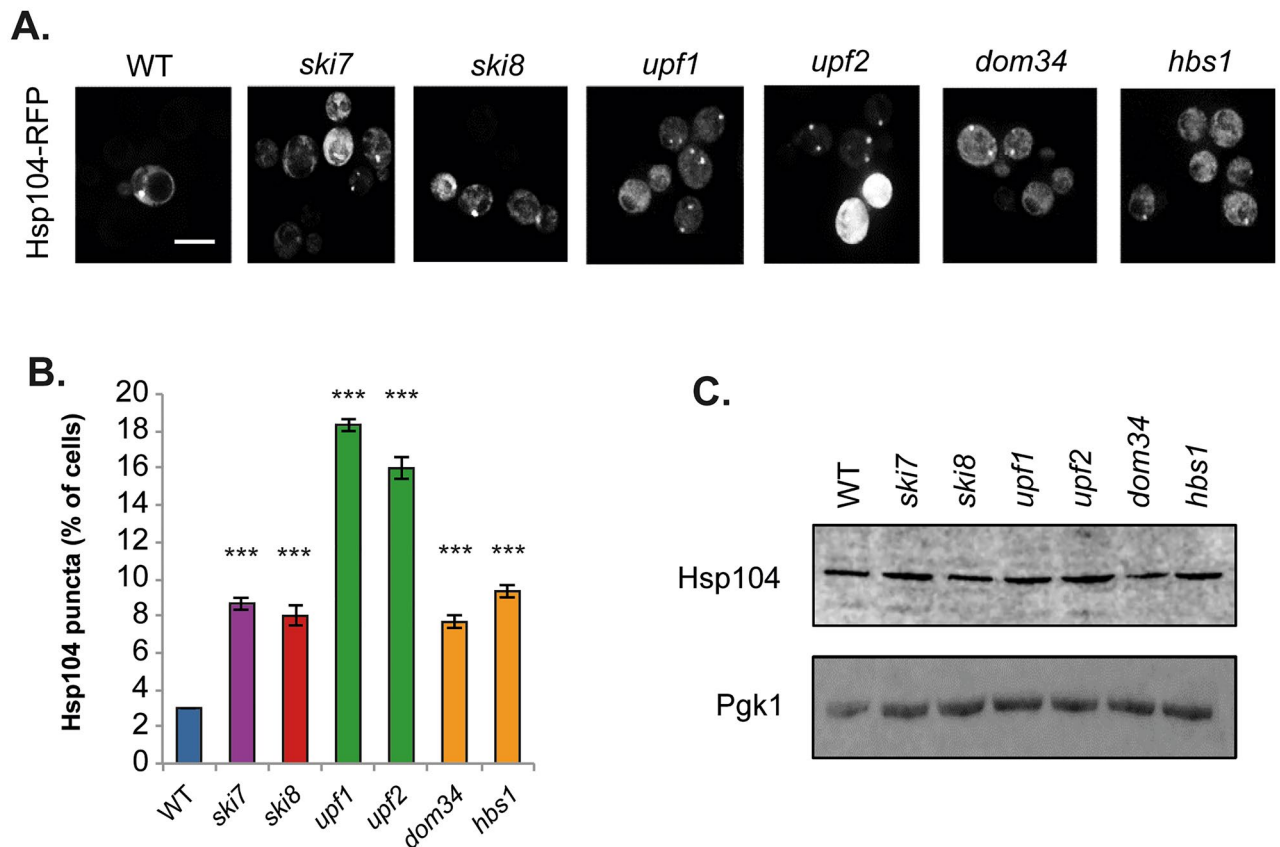



Figure 1. Strains lacking components of mRNA surveillance pathways have higher levels of protein aggregation. (A) Hsp104-RFP was visualized in wild-type and mutant strains disrupted for NGD (*dom34*, *hbs1*), NMD (*upf1*, *upf2*), NSD (*ski7*) and the Ski complex (*ski8*). Examples of cells containing visible puncta are shown. (B) The percentage of cells containing visible Hsp104-RFP puncta is quantified for each strain. Data shown are the means of three independent biological repeat experiments \pm SD. Significance is shown compared with the wild-type strain; *** $p < 0.001$. (C) Western blot analysis of Hsp104 protein levels. Blots were probed with a Pgk1 antibody as a loading control. The full blots are shown in Supplementary Fig. 1.

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