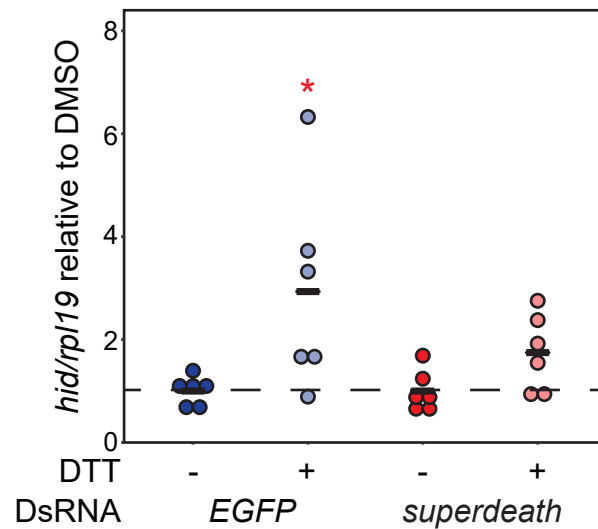


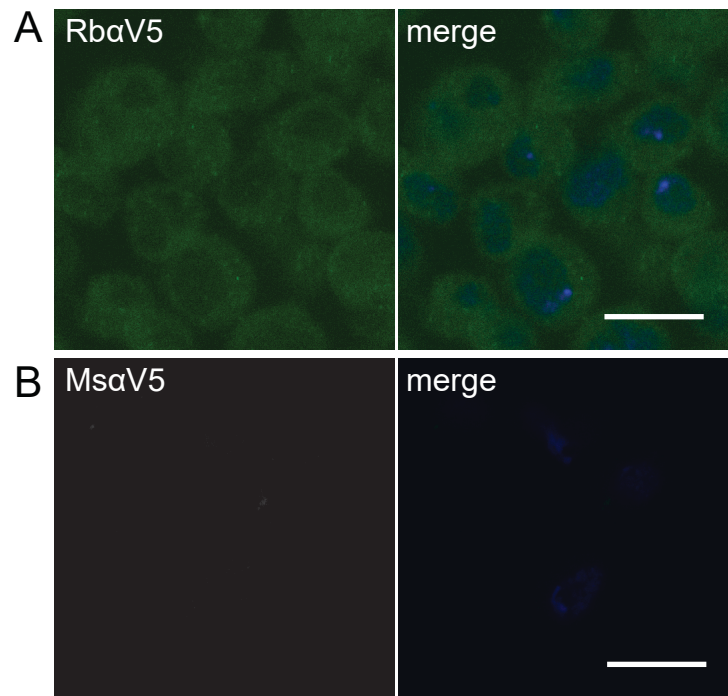
**Supplementary Figure 1. *superdeath* expression is efficiently reduced by both RNAi and DsRNA.**

The RNAi and DsRNA targeting *superdeath* in our models of ER stress significantly reduce *superdeath* expression. **A.** The Bloomington *Drosophila* Stock Center *superdeath* RNAi strain (42947) used in this study efficiently reduces expression of *superdeath* (27.8% of controls, N = 4). The RNAi construct was driven ubiquitously by *Tubulin-GAL4*, and expression determined in wandering L3 larvae compared to controls expressing only *Tubulin-GAL4* (N = 4). The control used for this RNAi line throughout the study is the *attP40* strain (Bloomington #36304). **B.** A second RNAi strain for *superdeath* (Bloomington #35802) causes a similar increase in eye size when *Rh1*<sup>G69D</sup> is overexpressed. Degeneration caused by overexpression of *Rh1*<sup>G69D</sup> is partially rescued by RNAi-mediated knockdown of *superdeath* expression (16391  $\pm$  1603 pixels, N = 36 in *Rh1*<sup>G69D</sup>/*superdeathi* flies as compared to 15174  $\pm$  1039 pixels, N = 33 in *Rh1*<sup>G69D</sup>/controls). Alternate control is the *attP2* strain (Bloomington #36303). **C.** A third RNAi strain for *superdeath* (Vienna #108616) causes a similar increase in eye size when *Rh1*<sup>G69D</sup> is overexpressed. Degeneration caused by overexpression of *Rh1*<sup>G69D</sup> is partially rescued by RNAi-mediated knockdown of *superdeath* expression (15403  $\pm$  1941 pixels, N = 21 in *Rh1*<sup>G69D</sup>/*superdeathi* flies as compared to 14371  $\pm$  1225 pixels, N = 52 in *Rh1*<sup>G69D</sup>/controls). Alternate control is the *attP* strain (VDRC #60100). **D.** S2 Cells treated with DsRNA targeted against *superdeath* also had near complete reduction in *superdeath* transcript levels (10.2%, N = 6) as compared to control cells treated with DsRNA against *EGFP* (N = 6). Values are average  $\pm$  SD. \* P < 0.05, \*\*\* P < 0.0005, \*\*\*\* P < 0.00005.



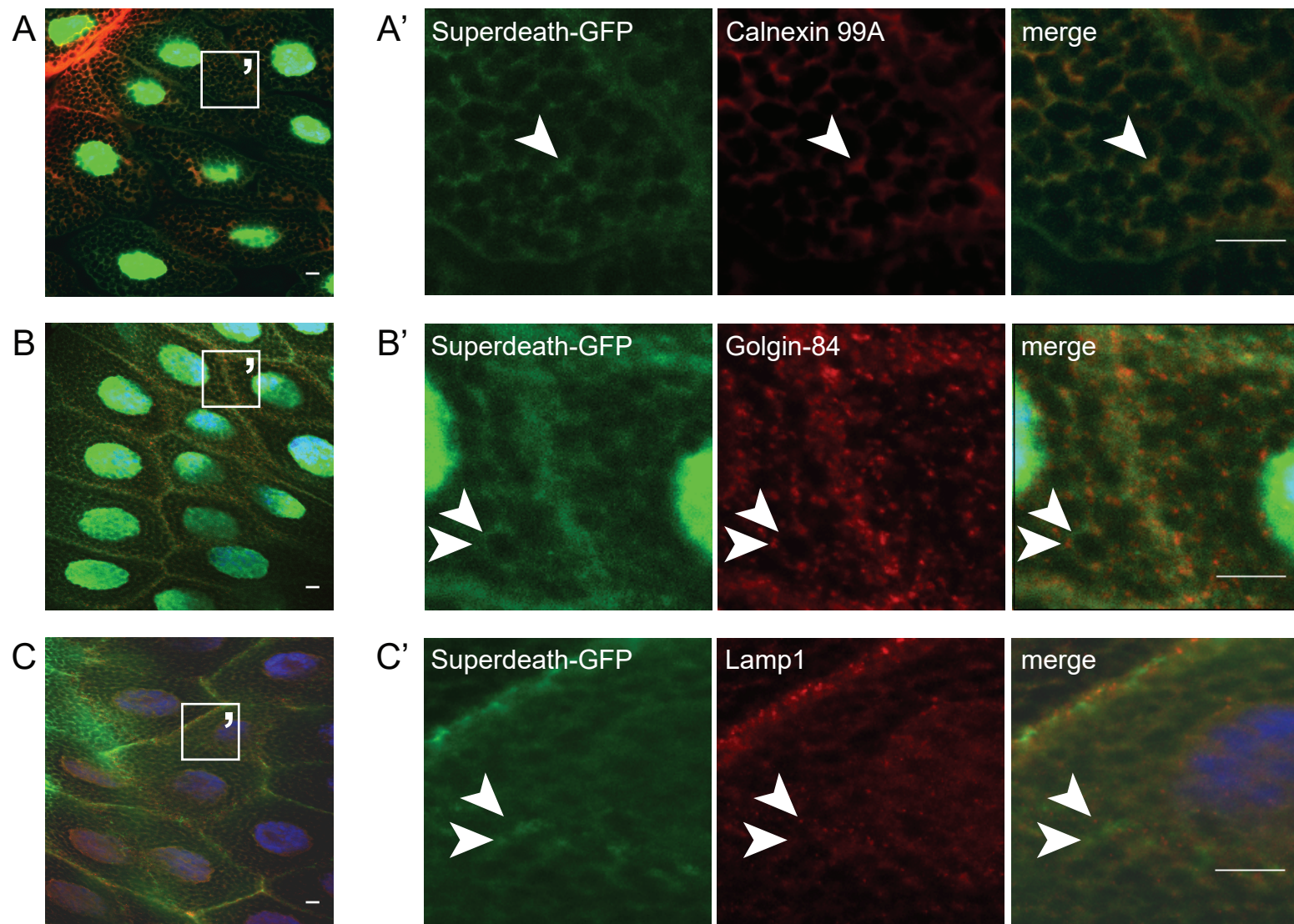
**Supplementary Figure 2. Loss of *superdeath* reduces the upregulation of apoptotic transcripts under ER stress.**

S2 cells treated with DsRNA against EGFP increased expression of the apoptotic gene *hid* after 7.5 hours of DTT exposure ( $2.93 \pm 1.98$ , N = 6) as compared to DMSO-treated control cells ( $1.00 \pm 0.284$ , N = 6). *hid* expression was not significantly increased in S2 cells that were treated with DsRNA against *superdeath* ( $1.74 \pm 0.75$ , N = 6 in DTT-treated cells compared to  $1.00 \pm 0.40$ , N = 6 with DMSO).



**Supplementary Figure 3. Specificity of V5 staining.**

The signal for Superdeath-V5 in Figure 6 is specific to expression of the V5 tag. A. S2 cells that were not transformed with the pMT-DEST48-Superdeath-V5 vector were stained for rabbit  $\alpha$ V5 (green) and counterstained with DAPI. Only background staining and no punctate staining indicative of Superdeath is detectable. B. S2 cells that were not transformed with the pMT-DEST48-Superdeath-V5 vector were stained for mouse  $\alpha$ V5 (green) and counterstained with DAPI. No punctate staining indicative of Superdeath is detectable. Scale bars = 0.01 mm.



**Supplementary Figure 4. Superdeath is localized to the endoplasmic reticulum *in vivo*.**

Superdeath predominantly localizes to the ER membrane in the salivary glands. **A.** Superdeath localizes to the ER. Salivary glands expressing Superdeath-GFP (Bloomington #64447) were stained for GFP (green, Rb antibody Thermo-Fisher #A6455) and Calnexin 99A (red) and counterstained with DAPI. **A'** represents the highlighted panel from **A**. The white arrow highlights select sites of GFP and Calnexin 99A overlap (orange). **B.** Superdeath does not primarily localize to the golgi. Salivary glands expressing Superdeath-GFP were stained for GFP (green, Rb antibody Thermo-Fisher #A6455) and Golgin-84 (red) and counterstained with DAPI. **B'** represents the highlighted panel from **B**. White arrows indicate select sites of independent GFP staining (green) or Golgin-84 staining (red). **C.** Superdeath does not primarily localize to the lysosome. Salivary glands expressing Superdeath-GFP were stained for GFP (green, Ms antibody MBL #M048-3) and Lamp1 (red) and counterstained with DAPI. **C'** represents the highlighted panel from **C**. White arrows indicate select sites of independent GFP staining (green) or Lamp1 staining (red). Non-specific nuclear staining is only seen in cells stained with the rabbit  $\alpha$ GFP antibody. Scale bars = 0.01 mm.