

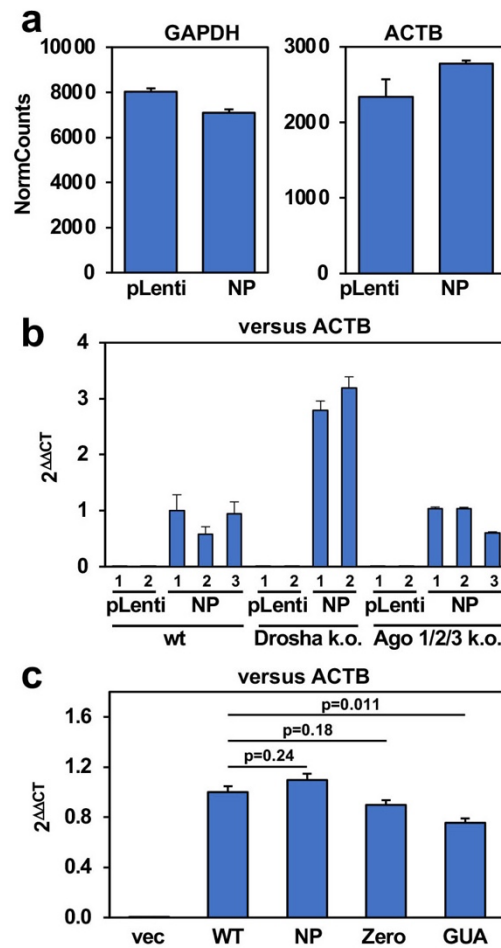


Figure S7 - Reanalysis of the qPCR data in Fig. 5b and 6b normalized to ACTB

(a) RNA-Seq analysis of GAPDH and ACTB expression in CD95 k.o. HeyA8 cells 50 hrs after infection with either pLenti or pLenti-CD95L (previous data deposited at GSE103631 were analyzed). Shown are the means of duplicate samples with SD. Raw read counts were normalized to 1 million reads per sample (NormCounts). (b) Reanalysis of the data shown in Fig. 5b normalized to ACTB, Bars represent the mean with standard deviation of triplicates. (c) Reanalysis of the data shown in Fig. 6b normalized to ACTB, Bars represent the mean with standard deviation of triplicates. T-test, Bonferroni adjusted p-value.