

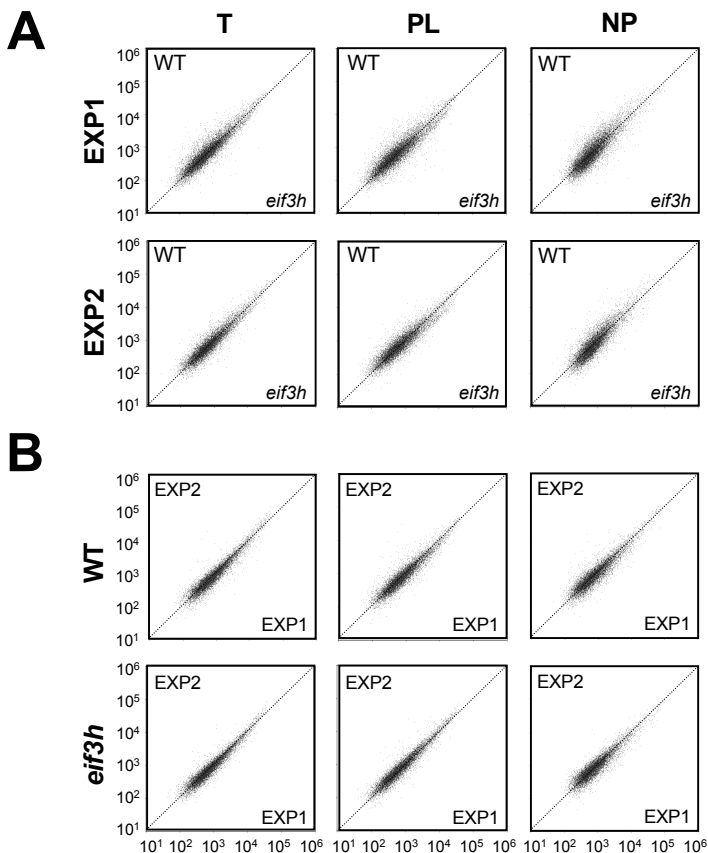


Figure S1. Scatter plots generated from microarray signal intensities after standard normalization. Genes with “absent” calls were excluded. **(A)** Comparison of signal intensities between wild type and *eif3h* mutant for total transcript (TC), polysome (PL) and non-polysome (NP) samples. **(B)** Comparison of signal intensities between two replicate experiments for total transcript levels (TC), polysomal (PL) and non-polysomal (NP) RNAs. The variance between experimental repeats (B) was smaller than that between wild-type and mutant samples (A), implying that the reproducibility between the two biological replicates was generally high.