**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.