



Supplemental Figure 4 Gene Ontology (GO) enrichment of overlapping long-lived mRNAs (LLRs).

(a, b) LLRs with decrease in transcripts level ($\log_2\text{FC} < 0$) in the embryo (a) and aleurone layer (b). (c, d) LLRs with significant increase in transcripts level ($\log_2\text{FC} \geq 1$, $\text{FDR} < 0.05$) in the embryo (c) and aleurone layer (d). The y-axis represents the functional subcategory, and the x-axis represents the ratio in the foreground list, which indicates the significance level and number in the foreground list.