



Supplemental Figure 1. Expression of *Lsi6* in different tissues normalized by Ubiquitin expression.

(A) Temporal variation of *Lsi6* expression in different rice tissues at the reproductive stages from before heading to grain filling. (B) Spatial expression of *Lsi6* in different rice tissues at flowering stage. The expression was determined by quantitative RT-PCR and expression level of Ubiquitin was used as an internal control. Expression relative to the root expression before heading (A) or unelongated stem (B) is shown. Data are means \pm SD of three biological replicates ($n = 3$).

