



Figure S4. Differential expression analysis of sncRNA and lincRNA

- a. The differential expression analysis of miRNA identified 686 up-regulated and 197 down-regulated miRNA during meiosis. The expression of miRNA in different meiotic stages was compared to flag leaf (control).
- b. The differential expression analysis of siRNA identified 121,341 up-regulated and 19,162 down-regulated siRNA during meiosis. The expression of siRNA in different meiotic stages was compared to flag leaf (control).
- c. The differential expression analysis of lincRNA identified 1,584 up-regulated and 217 down-regulated lincRNA during meiosis. The expression of lincRNA in different meiotic stages was compared to flag leaf (control).