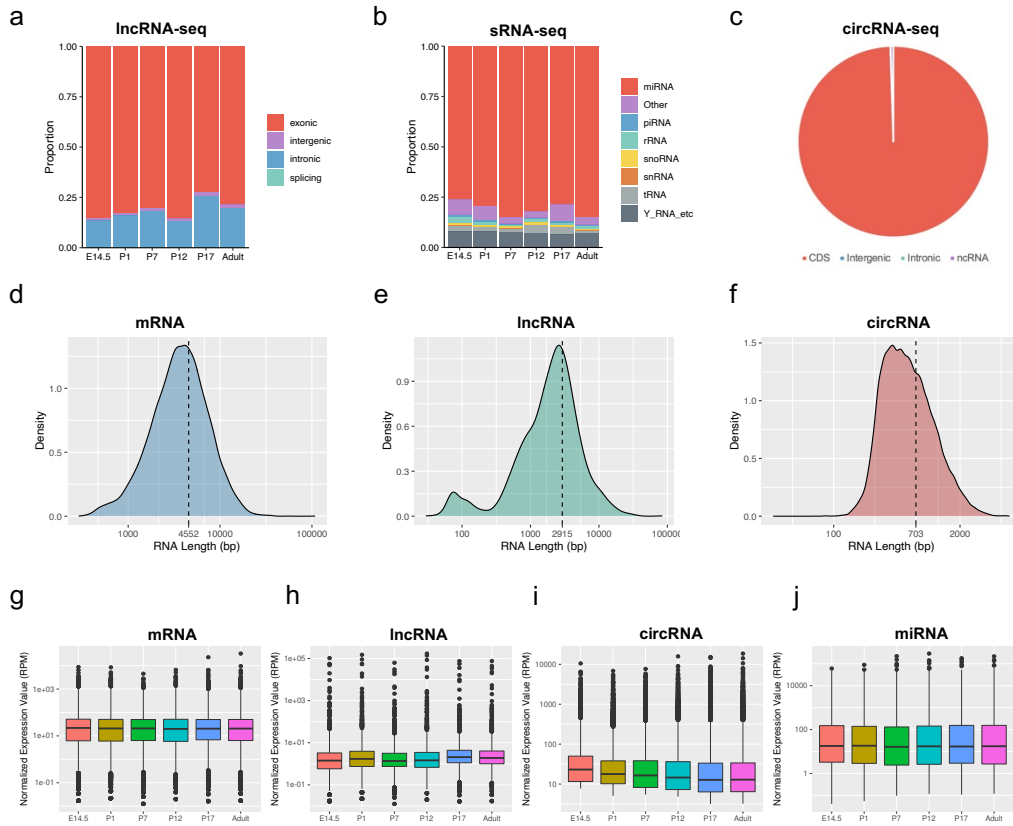
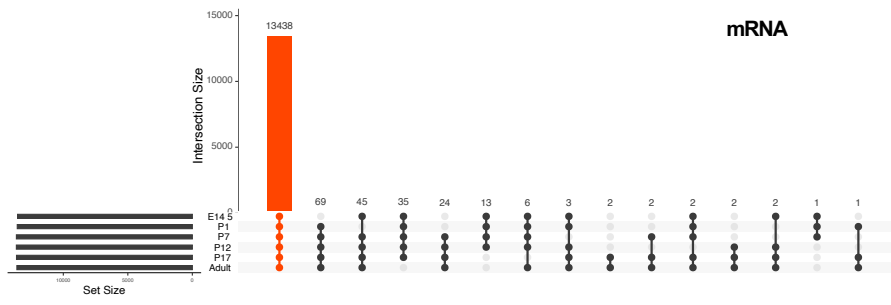


## Supplemental Figures

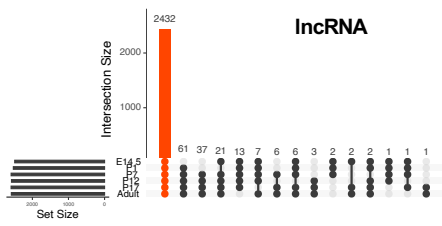


**Supplementary Figure 1. Whole-transcriptome sequencing data summary. (a-c)** Transcripts annotation from lncRNA-seq, sRNA-seq and circRNA-seq. **(d-f)** Length density plot of mRNA, lncRNA and circRNA. **(g-j)** Expression abundance of four class of transcriptomes across six developmental stages.

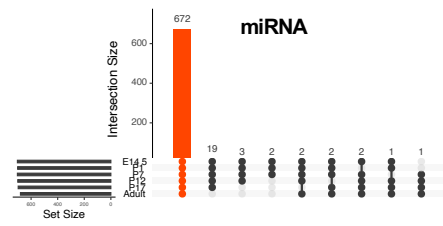
a



b



c



**Supplementary Figure 2. Upset plot of intersected transcripts in six stages. (a-c) Overlapped lncRNA, mRNA and miRNA numbers in different stages.**