

Additional data file 1: Supplementary Figures and Legends Jinming Huang *et al.*

Figure S1 The overall flow of the sequencing and informatic analysis for small RNA. (A) RNA sequencing experiment process of bovine tissue. (B) Bioinformatics of Solexa sequencing data.

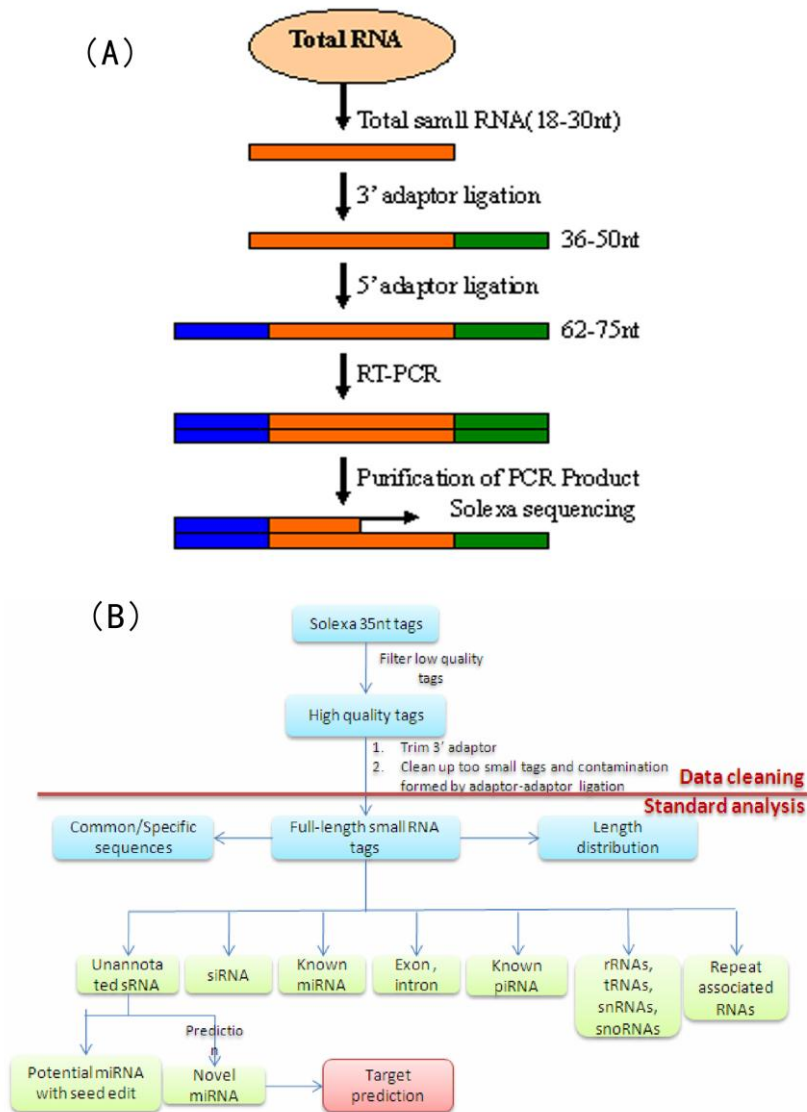


Figure S2. Summary of common and specific sequences between bovine testis (G1) and ovary (L4). (A) Number of unique small RNA tags. (B) Number of total small RNA tags.

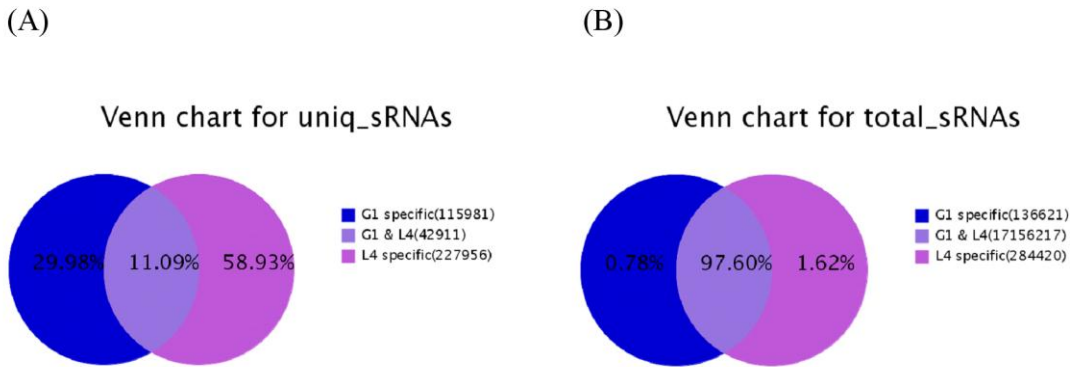


Figure S3. Distribution of small RNAs among different categories in testis (G1) and ovary (L4)

