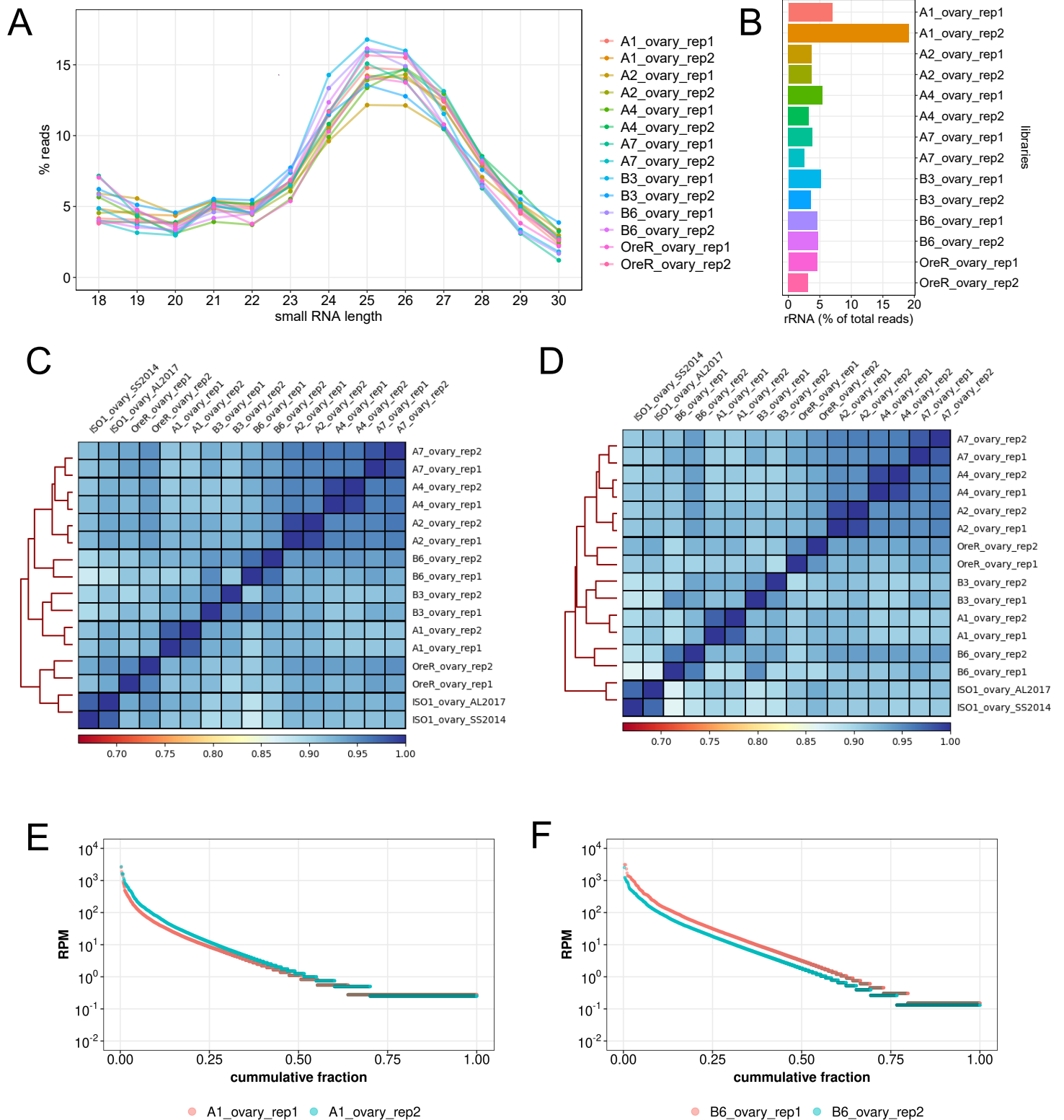


Supplementary Fig. 1



Supplementary_Fig_S1. Small RNA library QC results. (A) Small RNA read length profile after processing and removal of miRNA, tRNA, rRNA and snoRNAs. (B) Percentage of rRNA derived reads in small RNA libraries. (C&D) Spearman correlation plot of TE-derived small RNA expression in libraries using unique and multi mapping approaches respectively. (E&F) Comparison of cumulative sequence distribution of piRNAs in libraries to evaluate saturation of sequence space.