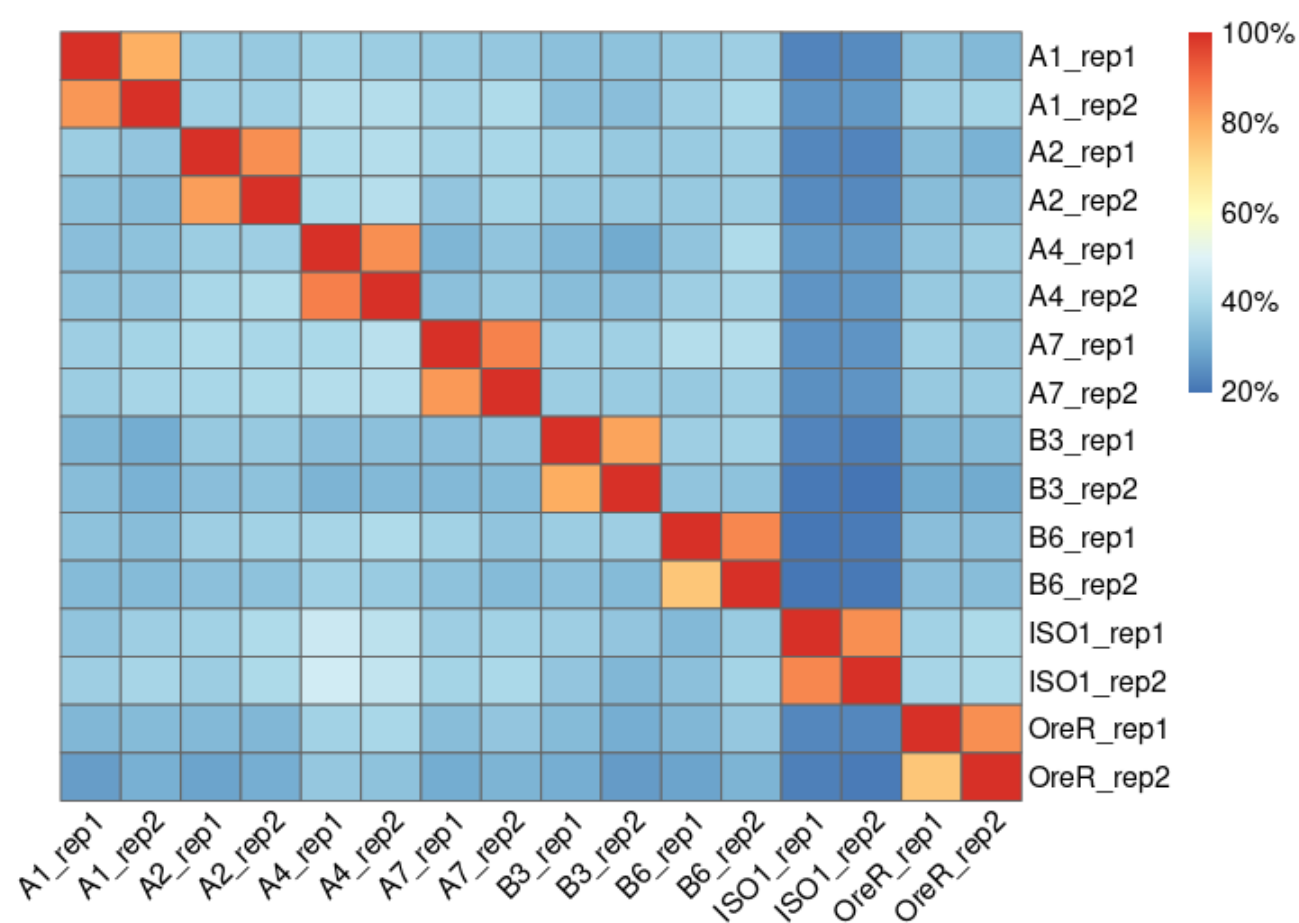
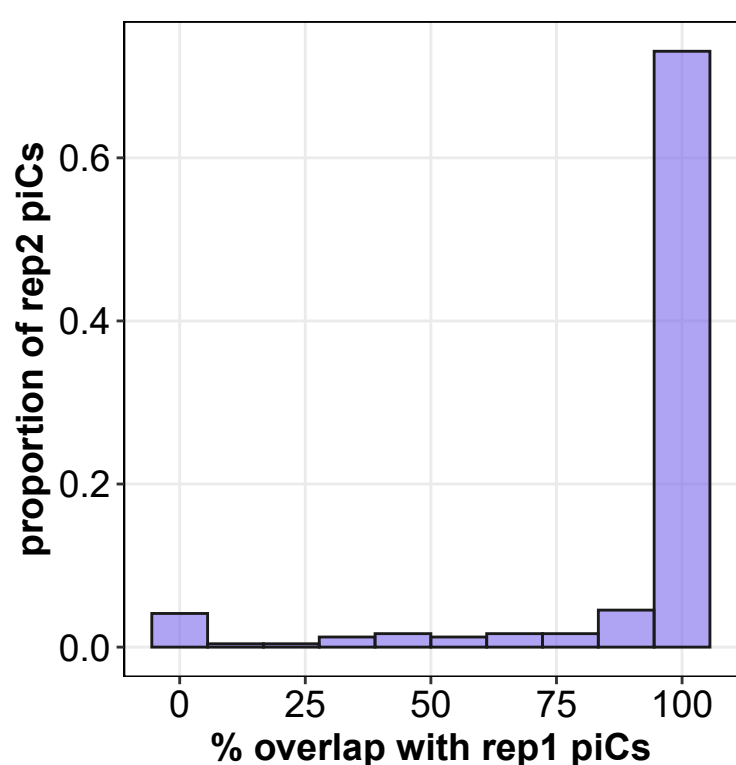


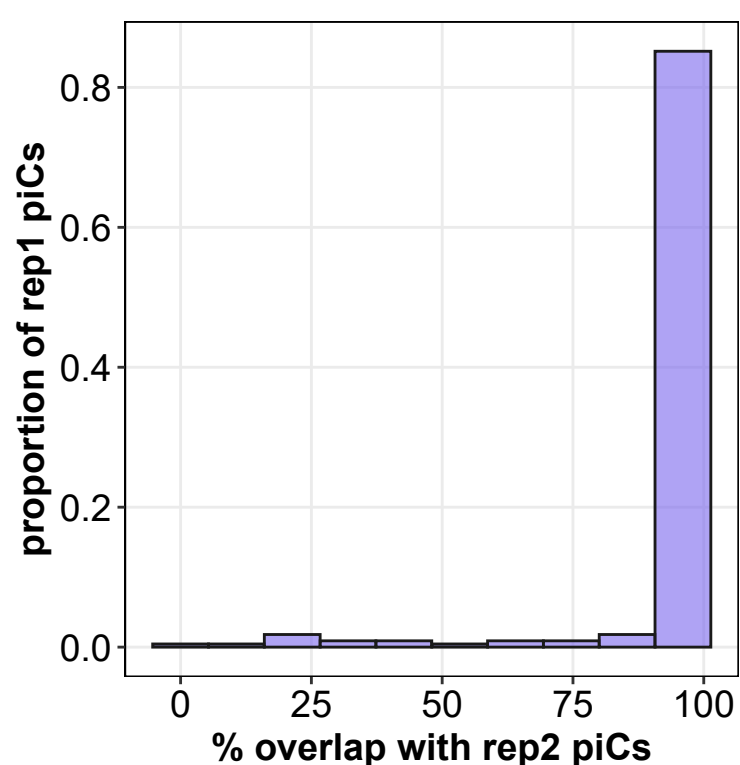
A



B



C



Supplementary_Figure_S3. Reproducibility of piC annotations. (A)) Cross strain and replicate overlap (% of total piCs count) of independently predicted piCs for each small RNA library using the restrictive method. Inter-strain comparisons are done after liftover to iso-1 coordinates and overlap >1bp between two piCs is deemed as shared. Intra-strain comparisons are done similarly but in native genome assembly. (B&C) Histogram showing proportion of piCs that overlap 0-100% of their respective length in annotations between two biological replicates of strain A1 replicate 1 and replicate 2 small RNA libraries.