

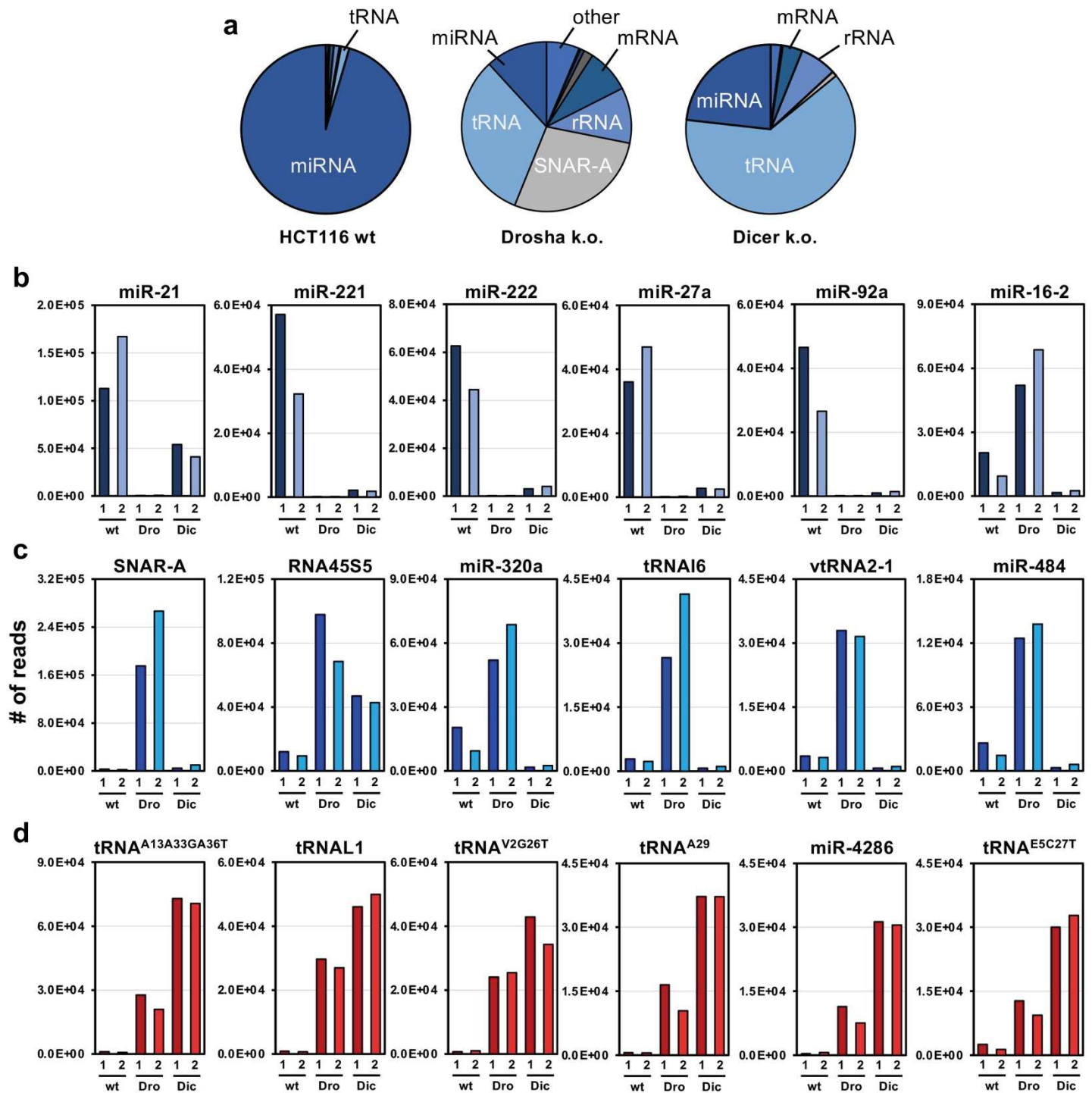


Figure S3 - Characterization of R-sRNAs in cells lacking Dicer and Drosha

(a) Pie charts representing the types of sRNAs that occupy the RISC of HCT116 cells by genotype. (b) Number of reads normalized per million of the top five most abundant sRNAs in the RISC of HCT116 cells (navy), (c) HCT116 Drosha k.o. cells (blue), and (d) HCT116 Dicer k.o. cells (red). Each bar represents read counts in one sample replicate, sample replicate 1 (darker shade) and replicate 2 (lighter shade).