Absolute distribution of small RNA sequences in three classes according to annotation.
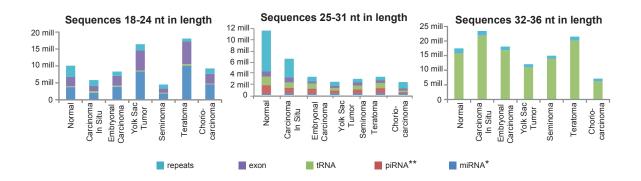The sequences are divided into three categories based on the peaks in the size distribution,
corresponding to miRNAs, piRNAs and longer sequences of previously unknown origin. The histograms
indicate the average sequence count per sample in each group.
* As annotated by miRBase, ** As annotated by piRNABank