



Supplemental Figure 1. Distribution of non-templated nucleotides as a percentage of all 4 nucleotides at both 5' and 3' positions relative to the mature miRNA or equivalent non-miRNA sequence. miRNA reads show a strong bias for non-templated A and U nucleotides in the 3' region, consistent with our general knowledge of isomiR structure. The non-miRNA reads had consistent nucleotide usage across the reads.