



Supplemental Figure S4: Statistics of high confidence exon junctions with GNR $\geq 2\%$.

(A): Distribution of gene expression (MRIG, \log_{10} scale) in 6,030 exon junctions with non-canonical dinucleotides.

(B): Average number of samples for all 191,448 exon junctions with GNR $>2\%$ (mean \pm SEM), grouped by dinucleotide.

(C): Average number of samples for all 191,448 exon junctions with GNR $>2\%$ MRIG (mean \pm SEM), grouped by GNR.

(D): Average number of samples for all 191,448 exon junctions with GNR $>2\%$ (mean \pm SEM), grouped by absolute number of reads (\log_{10} scale).