Figure S2. Variant allele frequency plots for assessing transcriptome-only mapping strategy
The variant allele frequencies of the SNVs that have at least 20X reads in exome-seq and RNA-seq are plotted. The RNA-seq SNVs were obtained with the transcriptome-only alignment option. Red and green dots represent the SNVs that are detected only in RNA-seq and only in exome-seq, respectively, while black dots represent the SNVs that are called in both. Venn diagrams are produced from the points represented in the graphs. The plots are generated for (A) RPMI8402 cell line and (B) TLE79 patient sample.