



Supplemental Fig S13. GO analysis of genes that harbor editing sites in the 3'ss (AG to GG). GO categories corresponding to different types of biological functions are color-coded (see Box). (A) Enriched GO terms among genes that contain Alu-overlapping exons with 3'ss editing. For this analysis, the background control genes consisted of those that have Alu-overlapping exons but without editing sites in their 3'ss. (B) Enriched GO terms among genes that contain non-Alu-overlapping exons with 3'ss editing. The background control genes were those that have non-Alu-overlapping exons but without editing sites in the 3'ss.