



S1 Fig. Proteomics screening of HuR related proteins. (A) The number of upregulated and downregulated proteins from the proteomics analysis after downregulation of HuR in TE-1. (B) KEGG pathway enrichment-based clustering analysis. (C) KEGG pathway-based enrichment analysis of upregulated proteins (shRNA-1 vs. shRNA-NC). (D) KEGG pathway-based enrichment analysis of downregulated proteins (shRNA-1 vs. shRNA-NC).