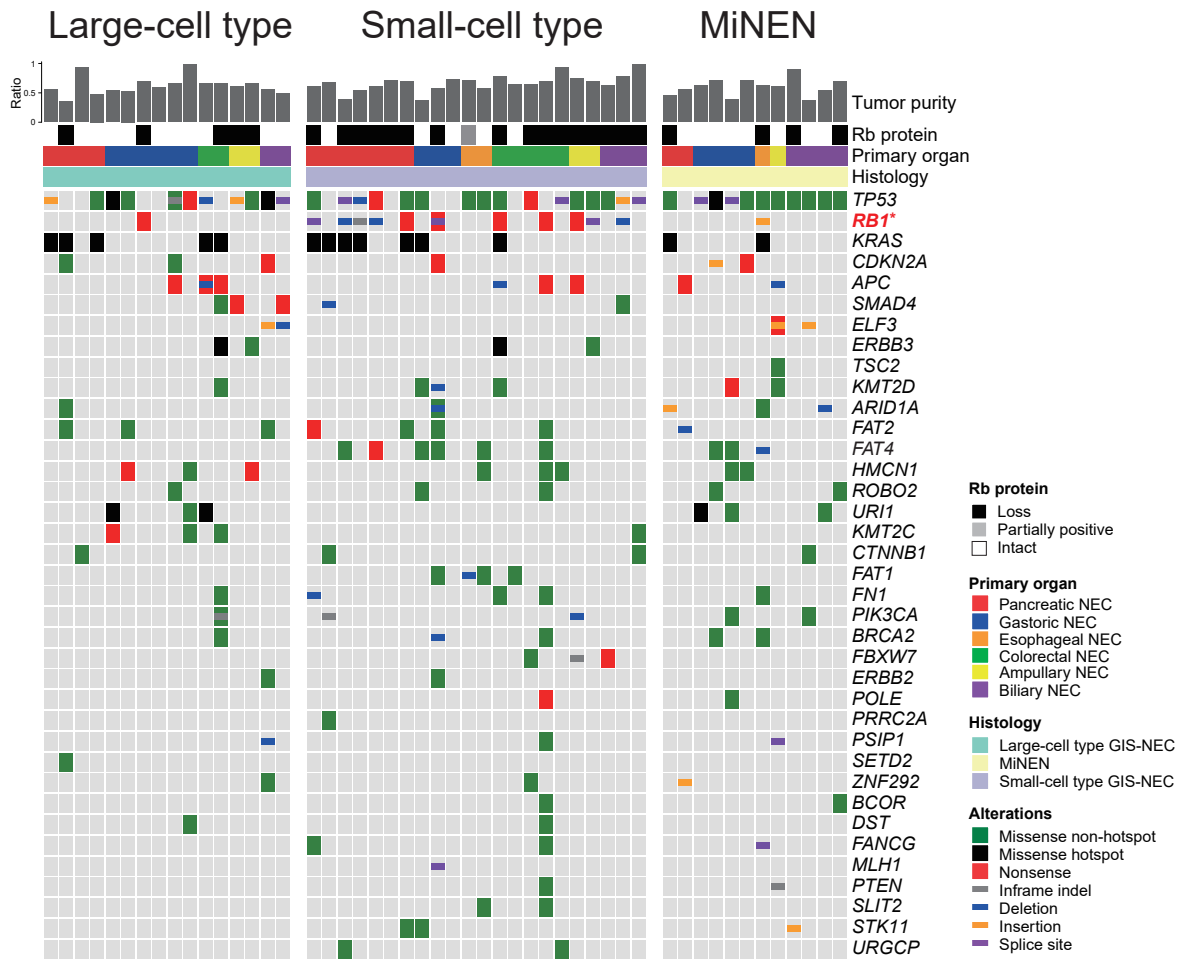
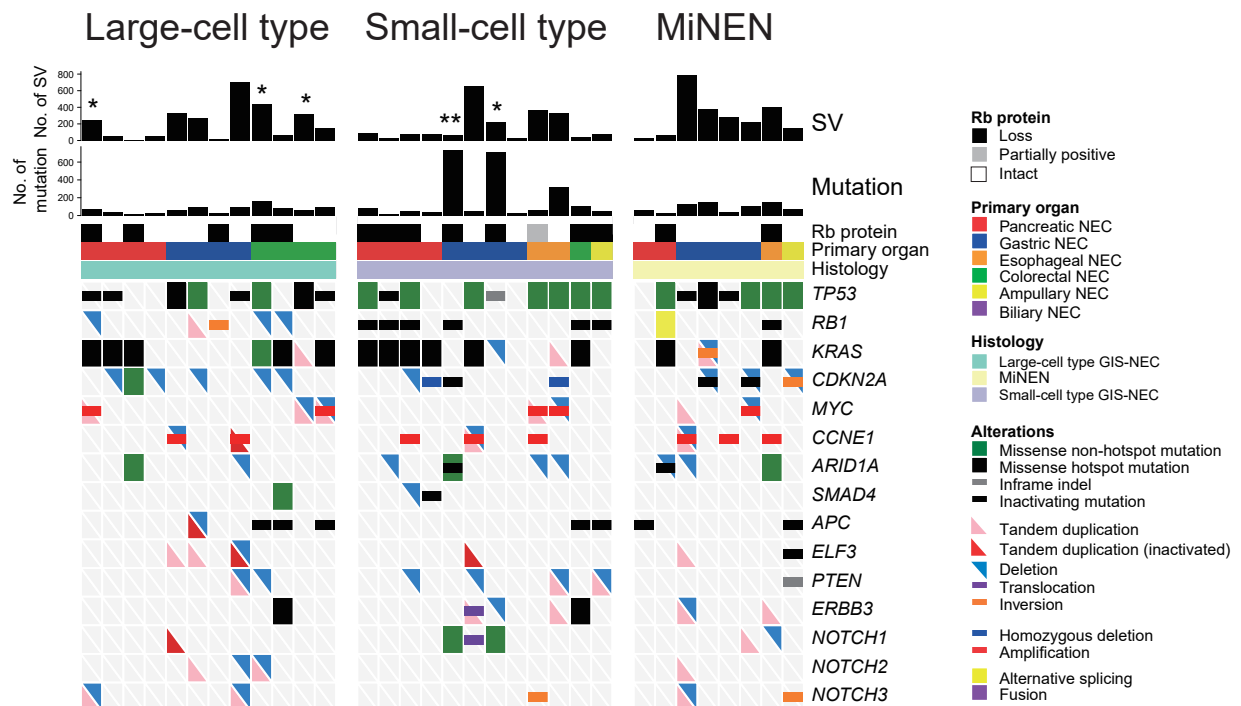


Supplementary Figure S7. Genomic landscape based on the morphological subclassification of GIS-NECs.

A



B



A, Oncoplot summary of mutated genes in large-cell type GIS-NECs, small-cell type GIS-NECs, and MiNEN based on targeted deep sequence analysis. The tumor purity, the status of Rb protein, and the primary organ are indicated at the top. **B**, Oncoplot summary of altered genes in large-cell type GIS-NECs, small-cell type GIS-NECs, and MiNEN based on WGS data. Data for SV and mutations, the Rb protein status, and the primary organ are provided at the top. *Asterisk*, organoid sample; *Double Asterisk*, hypermutator phenotype with somatic mutations of *MLH1* and *POLH* genes.