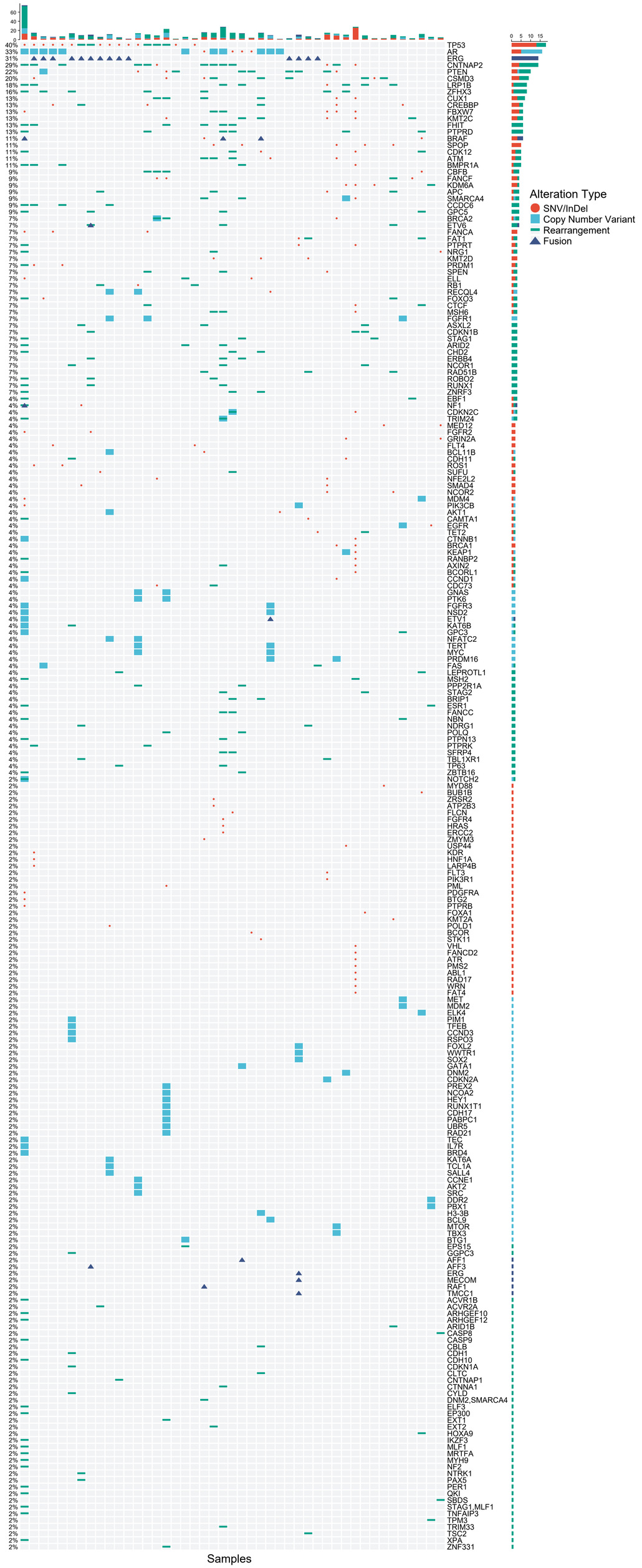

Supplementary Figure 1. Comprehensive landscape of genomic alterations detected by TE-WGS. Oncoprint summarizing all genomic alterations detected by TE-WGS across the cohort, including single nucleotide variants and indels, copy number variations (CNVs), deleterious rearrangements, and gene fusions. TE-WGS: Target-enhanced whole genome sequencing. SNV: Single nucleotide variant. Indel: Insertion/deletion. CNV: Copy number variation.