S7 Fig. Somatic mutation profiles of 35 SCLC patients from pre-treatment ctDNA (B1) sequencing of 1,021 genes. Thirty-five patients were arranged along the x-axis. Tumor mutation burden (muts/Mb) was shown in the upper panel. Genes with somatic mutations were shown in the middle panel. Mutation frequencies of each gene were shown on the left. Clinical characteristics for each patient was shown in the bottom.