Supplementary Figure 3. The immune contexture and genomic alteration based on NKG2A⁺CD8⁺ T cells infiltration. (A) Distribution of various immune cells in different infiltration groups of NKG2A⁺CD8⁺ T cells in ZSHC cohort; Green boxplots represent patients with low infiltration of NKG2A⁺CD8⁺ T cells, while red boxplots represent patients with high infiltration. Mann-Whitney U test. (B) The landscape of genomic alterations according to NKG2A⁺CD8⁺ T cells infiltration in TCGA-KIRC. Chi-square test. *P<0.05, **P<0.01, ***P<0.001, ns refers to no significance.