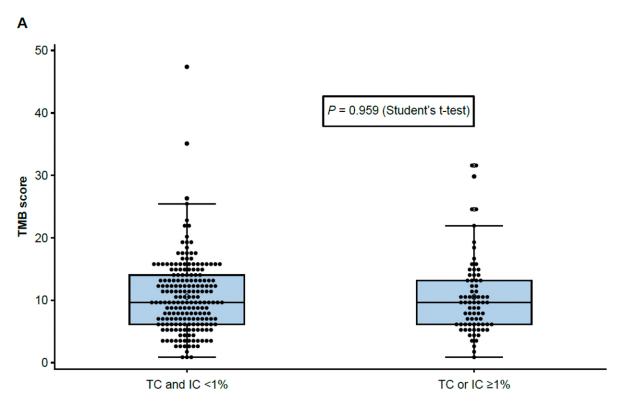
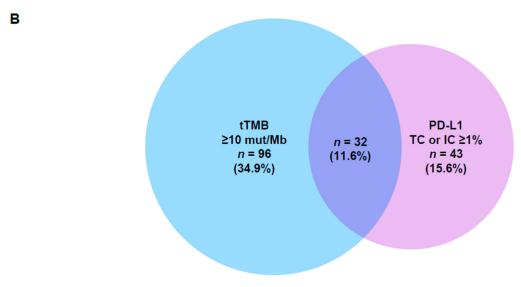
Supplementary Figure S11. Relationship between PD-L1 expression and tTMB score. (A) Box plots showing the distribution of tTMB score in patients by PD-L1 subgroup; (B) Venn diagram showing overlap between the tTMB ≥10 mut/Mb and PD-L1 TC or IC ≥1% subgroups. Percentages are calculated from the PD-L1 and tTMB BEP, i.e., patients who had both a PD-L1 and a tTMB result (n=275); samples from 171 patients had tTMB ≥10 mut/Mb and/or PD-L1 TC or IC ≥1%, while samples from 104 patients (37.8%) had both tTMB <10 mut/Mb and PD-L1 TC and IC <1%. BEP, biomarker evaluable population; IC, immune cell; mut/Mb, mutation per megabase; PD-L1, programmed cell death ligand-1; TC, tumor cell; TMB, tumor mutational burden.



PD-L1 expression



PD-L1 and tTMB BEP (n = 275)