

Figure S1. Tumor mutation profiles included high-frequency alterations in genes associated with (A) PFS and (B) OS. When tTMB was calculated, patients with tTMB-H had (C) longer PFS and (D) OS than patients with tTMB-L, although no significant differences were observed. mPFS, median progression-free survival; mOS, median overall survival; HR, hazard ratio; tTMB-H, tissue tumor mutation burden high.

