**Supplementary Fig. 2. Distribution of IGSHCC scores in HCC patients.** (A) TCGA training data set, (B) TCGA validation set, (C) entire TCGA data set, and (D) LCI validation data set. In each figure, the top panel represents the low- and high-IGSHCC groups defined based on the cutoff value for TCGA training data set, the middle panel represents the survival statuses and durations in the HCC patients in the low- and high-IGSHCC groups, and the bottom panel is a heatmap of the expression of the 22 immune-related genes in IGSHCC in each tumor.

