Supplementary Figure 1. Expression profiles of 57 hub genes with degree ≥20 in LUAD (A) and LUSC (B).
Supplementary Figure 2. Kaplan-Meier survival analysis for estimating the prognostic capacity of the ACK1 in TCGA lung cohorts. Risk group stratification by the ACK1 expression levels with respect to the overall survival of LUAD (A) and LUSC (B). Risk group stratification by the ACK1 expression levels with respect to disease-free survival of LUAD (C) and LUSC (D).