Fig. S2 Network analysis and functional enrichment analysis of SFKs

**Figure S1. Network analysis and functional enrichment analysis of SFKs.** (A) Protein-protein interaction network of individual SFKs (STRING). (B) Gene-gene interaction network of different SFKs (GeneMANIA). (C) GO functional enrichment on SFKs and the 20 associated genes, five significant entries from each part of biological process, cellular component and molecular function were displayed on the bubble chart (XIANTAO platform). (D) KEGG pathway analysis on SFKs and the related genes, 15 meaningful items were presented (XIANTAO platform). *PTK6*, *SRMS*, *YES1*, namely *BRK*, *SRM*, *YES*.