Supplementary Figure 1. The abundance of tumor infiltrating immune cells in the Cluster 1 and 2 estimated by TIMER2.0. TIMER2.0 estimated 6 kinds of tumor infiltrating immune cells.
Supplementary Figure 2. The abundance of tumor infiltrating immune cells in the Cluster 1 and 2 estimated by CIBERSORTx. CIBERSORTx estimated 22 kinds of tumor infiltrating immune cells.
Supplementary Figure 3. Protein-protein interaction network of 8 genes in the TME signature. TNFRSF1B and BLK with the most neighbor genes were identified as hub genes.