

Figure S1. Profiles of infiltrating immune cells in normal and PRCC tissues using GEO data. (a) Heatmap of 22 immune cell proportions in normal tissues, type 1 and type 2 PRCC tissues. (b) Violin plot showed the different expression of 22 immune cells between normal (blue), type 1 (red) and type 2 PRCC groups (green). *P < 0.05, **P < 0.01, ***P < 0.001.

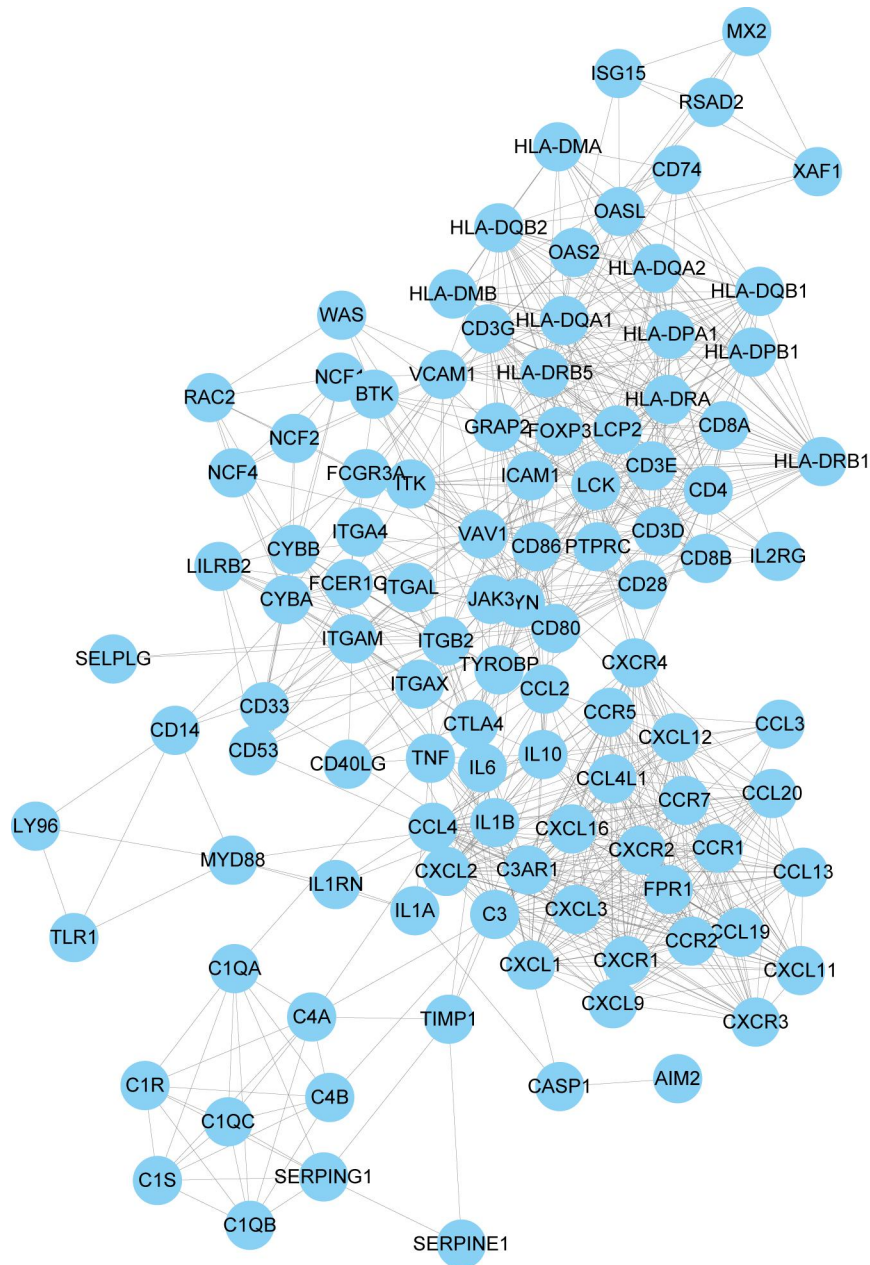


Figure S2. Protein–protein interaction network of differentially expressed genes in type 2 PRCC tissues.