

Figure S1 PPI network analysis of two module genes identified by WGCNA. The nodes indicate proteins; the lines represent the interaction of proteins. One thousand two hundred and fifty-four nodes and 7,754 edges were obtained with a confidence score of ≥ 0.7 . PPI, protein-protein interaction; WGCNA, weighted gene co-expression network analysis.

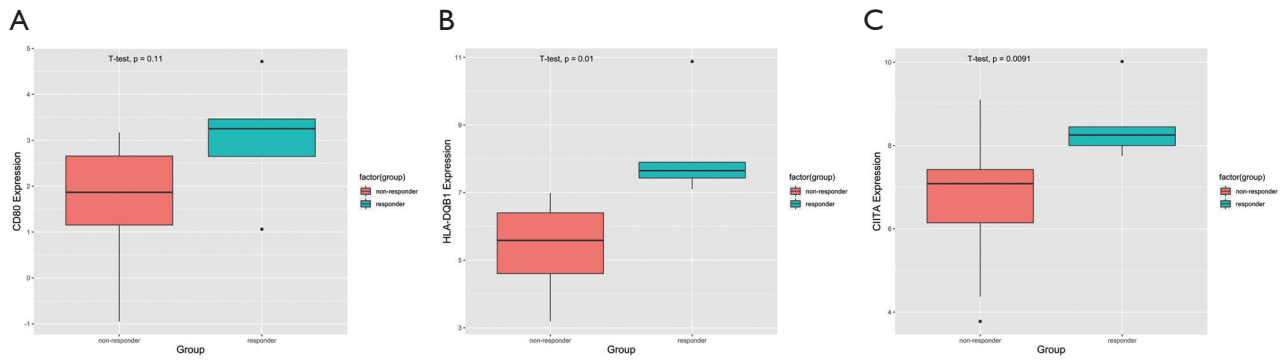


Figure S2 The expression of 3 DEmRNAs between the response group and non-response group based on GSE126044. (A) CD80; (B) HLA-DQB1; (C) CIITA. DEmRNAs, differentially expressed mRNA.