

SUPPLEMENTARY TABLE

Supplementary Table 1. Genes associated with good prognoses identified in TCGA.

Interferon-γ-mediated signaling pathway	CIITA, ICAM1, HLA-DQB2, HLA-A, HLA-C, OAS1, HLA-B, OAS2, HLA-E, TRIM22, HLA-DQA2, HLA-F, B2M, OASL, CD44, IRF5, IFNG, IRF8, IRF1, HLA-DPA1, HLA-DPB1, GBP2, GBP1, HLA-DRA
Antigen processing and presentation via MHC class II	HLA-DQB2, HLA-DPA1, HLA-DPB1, HLA-DMA, HLA-DQA2, HLA-DOB, HLA-DRA
Antigen processing and presentation via MHC class I	TAP1, HLA-A, HLA-C, HLA-B, HLA-E, TAPBP, HLA-F, B2M
Macrophage chemotaxis	CCL2, LGALS3, CCL5
Neutrophil chemotaxis	CCL2, LGALS3, S100A9, CCL8, CCL5, VAV1, CCL18, CCL17, CCL22, CCL13, CCL23, IFNG, CSF3R, XCL1, XCL2, SYK
Dendritic cell chemotaxis	CCR7, CCR5, CCR2, CCL19, CCL5
Positive regulation of T cell chemotaxis	CCR2, TNFSF14, CCL5, XCL1
B cell activation	IKZF3, CXCR5, LAX1, ZAP70, CD79A, CD40, PRKCB, BTK