

SUPPLEMENTARY TABLES

Supplementary Table 1. Differentially expressed immunity-related genes identified between breast cancer and normal tissues.

Regulation	Gene symbol
Up-regulated (69)	<i>CST4, CGA, VGF, CCL11, SCT, FABP6, S100A7A, TUBB3, CCL7, CCR8, IGHE, CAMP, PRLH, ORMI, TFR2, ULBP1, EPO, TNFRSF9, UMODL1, S100P, PROC, ARTN, OPRDI, IL21R, AMH, CXCL11, BIRC5, BMP8A, IL20, IL9R, CD1A, ORM2, ESM1, INHBA, PCSK1, PYY, IL27, TNFRSF18, BMPR1B, LTA, TRH, CCL20, ICOS, TMPRSS6, KCNH2, TNFSF4, EDN2, IL11, CCR4, ADM2, PDCD1, LCN12, ANGPTL6, WFIKKNI, CD19, OLR1, CXCL10, SCG2, SLC11A1, MMP12, TRAV26-1, OASL, PIK3R2, FGFR3, THPO, ULBP2, RBP2, TRAV39, INHA</i>
Down-regulated (110)	<i>ANGPTL7, ACVR1C, DEFB132, LEP, TSLP, LALBA, PENK, GDF10, RNASE7, EDN3, GFAP, RXRG, CCL14, SSTR1, PAK3, ADCYAPIR1, GPR17, CCL24, NTF4, NRG2, CXCL2, CMA1, ADIPOQ, AVPR2, TACR1, KL, PLXNA4, IL17B, BMP5, ANGPTL1, RBP4, OXTR, PROK1, NRG1, ANGPT1, DES, CXCL3, FAM3D, ALB, IGF1, ADRB2, PTGFR, FGF2, LEPR, SEMA3D, GREM2, BMP2, PTGS2, SAA2, S100B, CNTFR, IL6, TAC1, NTS, CD209, NPR1, CCL23, ADRB1, PTH1R, IL22RA2, PPARG, NR3C2, IL33, FABP4, PI15, PTX3, ROBO3, SEMA3A, SEMA6D, PI3, NTF3, GHR, FABP7, BMP6, TNFRSF10D, OGN, CTSG, LGR6, EGFR, LIFR, NR4A3, CCL13, SCGB3A1, SPINK5, CXCL5, MASP2, EDNRB, CCL28, IL17D, TGFBR3, SHC3, NPR3, NGFR, FGF10, FGF7, CXCL6, MIA, PTH2R, BACH2, NR5A2, FGF16, SEMA3G, FGF1, GNAI1, MARCO, CCL21, DEFB1, TNFRSF8, SAA1, S100A12</i>

Supplementary Table 3. Correlation of risk score with tumor stage in breast cancer.

Tumor stage	Number of patients	Risk score	P
i-iiia	497	1.205 ± 1.011	0.044
iib-iv	466	1.338 ± 1.054	