Figure S1. Association of *STAT3* expression with that of *Bcl-2*, *Bcl-xL* (*BCL2L1*), *Cyclin D1* (*CCND1*), *VEGF*, *matrix metalloproteinase* (*MMP*)*-2*, and *MMP-9* was determined from the breast cancer RNA-Seq data from The Cancer Genome Atlas Pearson test was used to determine the correlation coefficient and statistical significance.