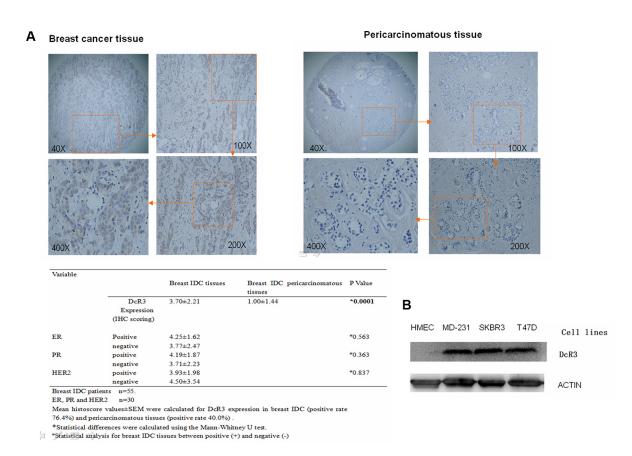
DcR3 gene polymorphisms are associated with sporadic breast infiltrating ductal carcinoma in Northeast Chinese women

Supplementary Materials



Supplemental Figure S1: Evaluation of DcR3 expression in breast cancer. DcR3 was detected in paraffin-embedded tissue sections using IHC analysis (A). IHC staining scores are reported as means \pm SE. The Mann-Whitney U test was used to analyze DcR3 expression differences between IDC and pericarcinomatous (PC) tissues $(3.70 \pm 2.21 \text{ to } 1.00 \pm 1.44)$. P = 0.0001. DcR3 was detected in HMEC and breast cancer cell lines via western blotting (B). DcR3 is overexpressed in breast cancer cell lines as compared to HMECs. Experiments were repeated 4 times.