

Supplementary Figure S3

Box and Whiskers plot showing the distribution of log2 transformed total TFPI ($\alpha+\beta$), TFPI α , and TFPI β and *TF* tumor mRNA expression across the following PAM50 subtype signatures of 150* breast cancer patients; basal (n=304/357), HER2 enriched (n=240/152), luminal A (n=465/482), luminal B (n=471/289) and normal-like (n=304/257). Data were derived from the GOBO database. *P*-values for ANOVA testing are indicated.

