Figure S2: Survival analysis in extreme quartiles of RERG and RLN2 mRNA expression in breast cancer data sets

Survival analysis in extreme quartiles of RERG and RLN2 mRNA expression in breast cancer data sets. RERG and RLN2 mRNA expression was determined in the indicated data sets as described in material and methods section and analyzed in Kaplan-Meier estimations. P-values are calculated based on log-rank tests.