



Figure S4. Kaplan-Meier plot showing the predictive power of the NR gene signature in an additional set of adenocarcinomas.

The NR gene-expression signature developed from the QPCR dataset (MDACC cohort, n=30) was validated in the microarray data from a cohort of 117 independent adenocarcinomas from Tomida et al. (18). *P*-values were obtained by the log-rank test. Red and black lines represent predicted high- and low-risk groups, respectively. Open circles indicate censored samples.