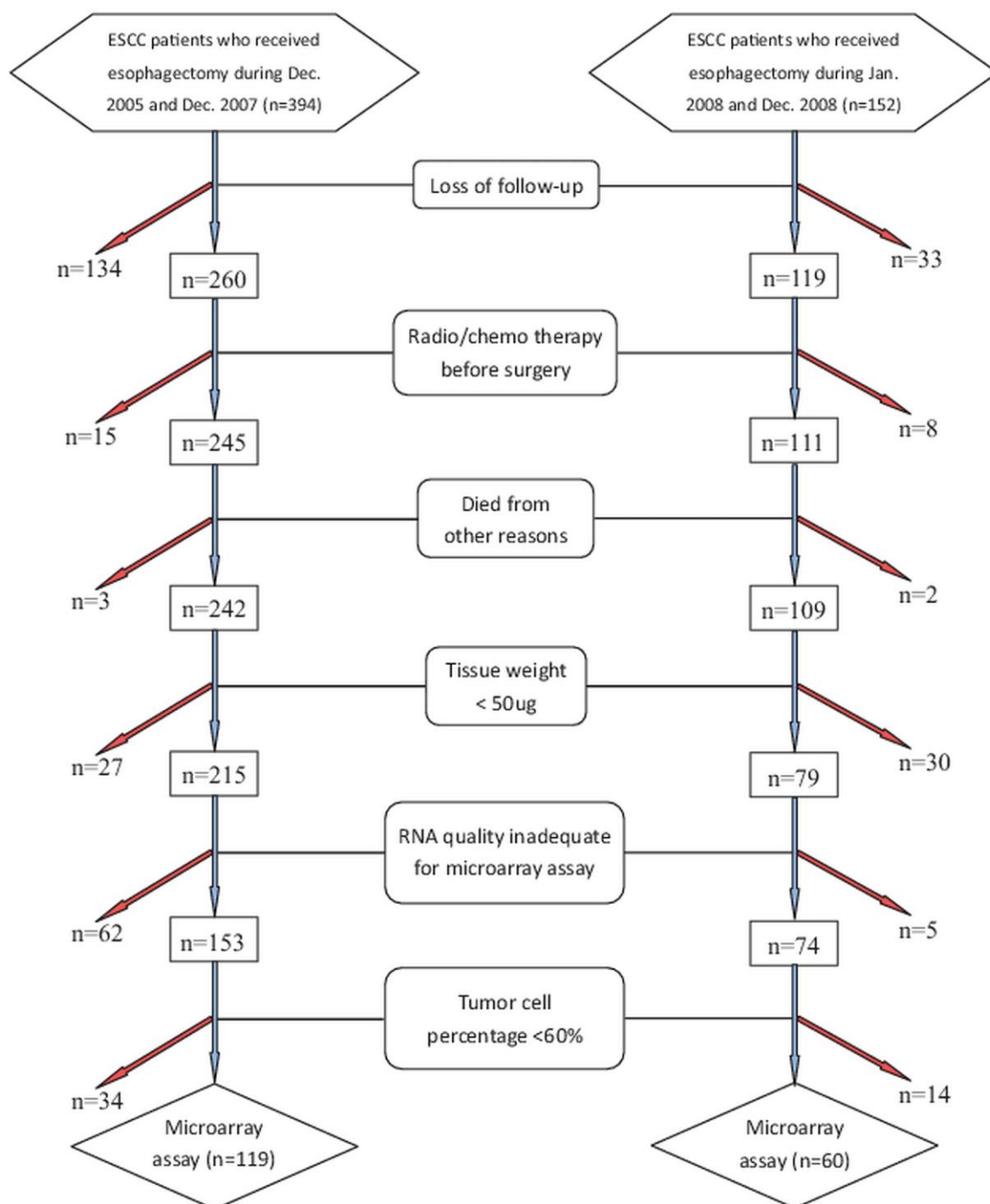


Supplementary Figure 1. Patient enrollment procedure diagram of the original (left) and independent (right) cohorts

Supplementary Figure 2. Comparison of lncRNA expression value measured by microarray and qRT-PCR.

The expression value of the three signature lncRNAs were measured by qRT-PCR and compared with microarray results in 25 randomly selected tumor samples. The quantitative RT-PCR reactions were performed in triplicate and the mean values of the lncRNAs were used. The expression levels of the 25 samples by microarray and qRT-PCR were shown for each lncRNA. For comparison, the expression value measured by both microarray and RT-PCR were normalized by z-score, shown as the longitudinal axis.

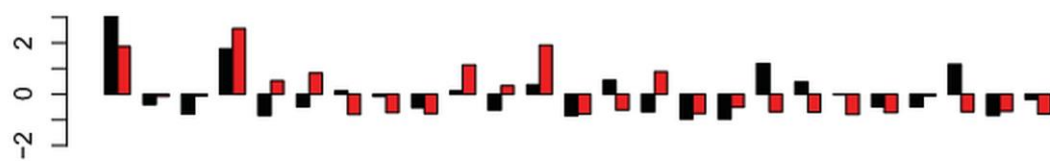
Supplementary Figures



ENST00000435885.1



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