Figure S1 All differentially expressed genes in module analysis via Cytoscape. Each blue square represents a gene, and the connected lines represent interactions between genes.
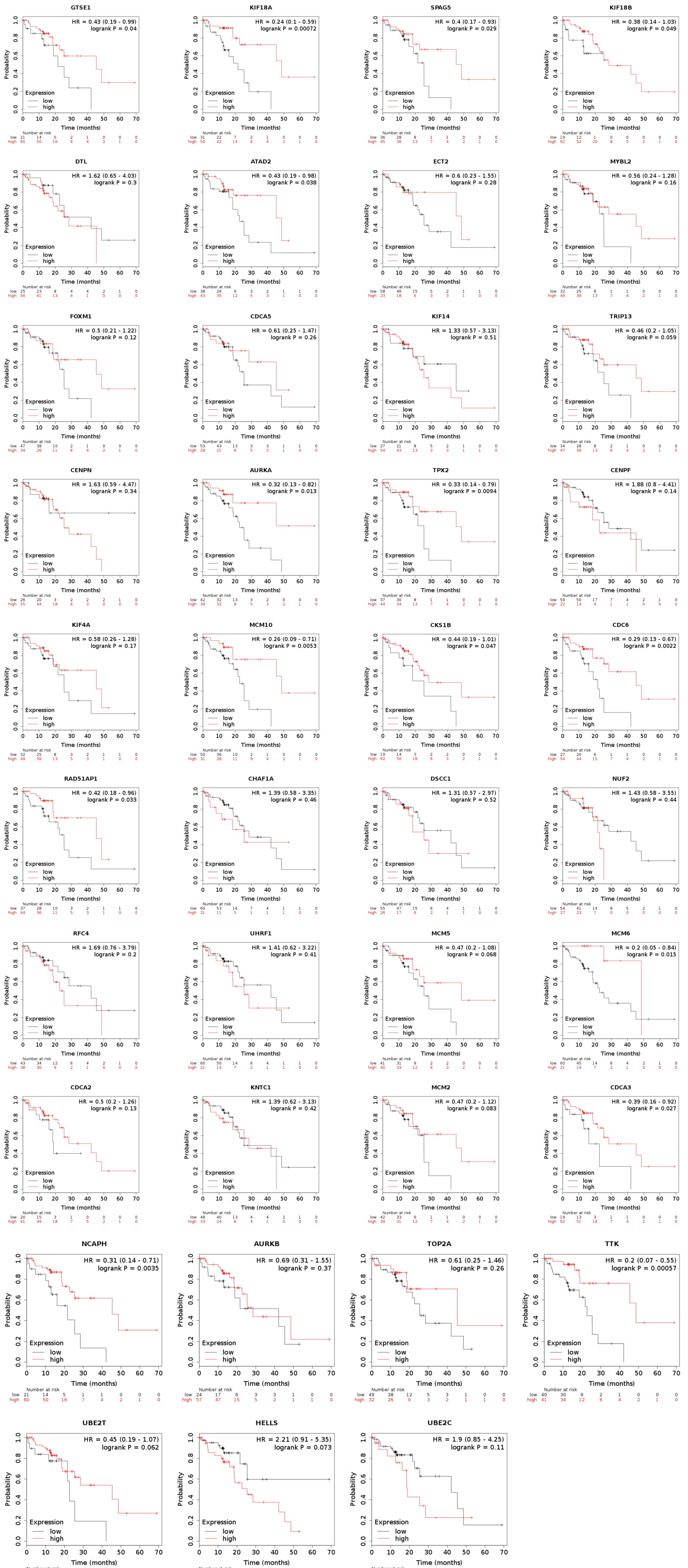
Figure S2 Survival curves of 39 hub genes (PICALM interacting mitotic regulator was excluded due to a lack of gene expression and survival information). Gene name is shown at the top of each figure part. Black line represents low expression of the identified gene, and red line represents high expression of the identified gene. Corresponding life table, hazard ratio, and log-rank P value are shown in each figure part.