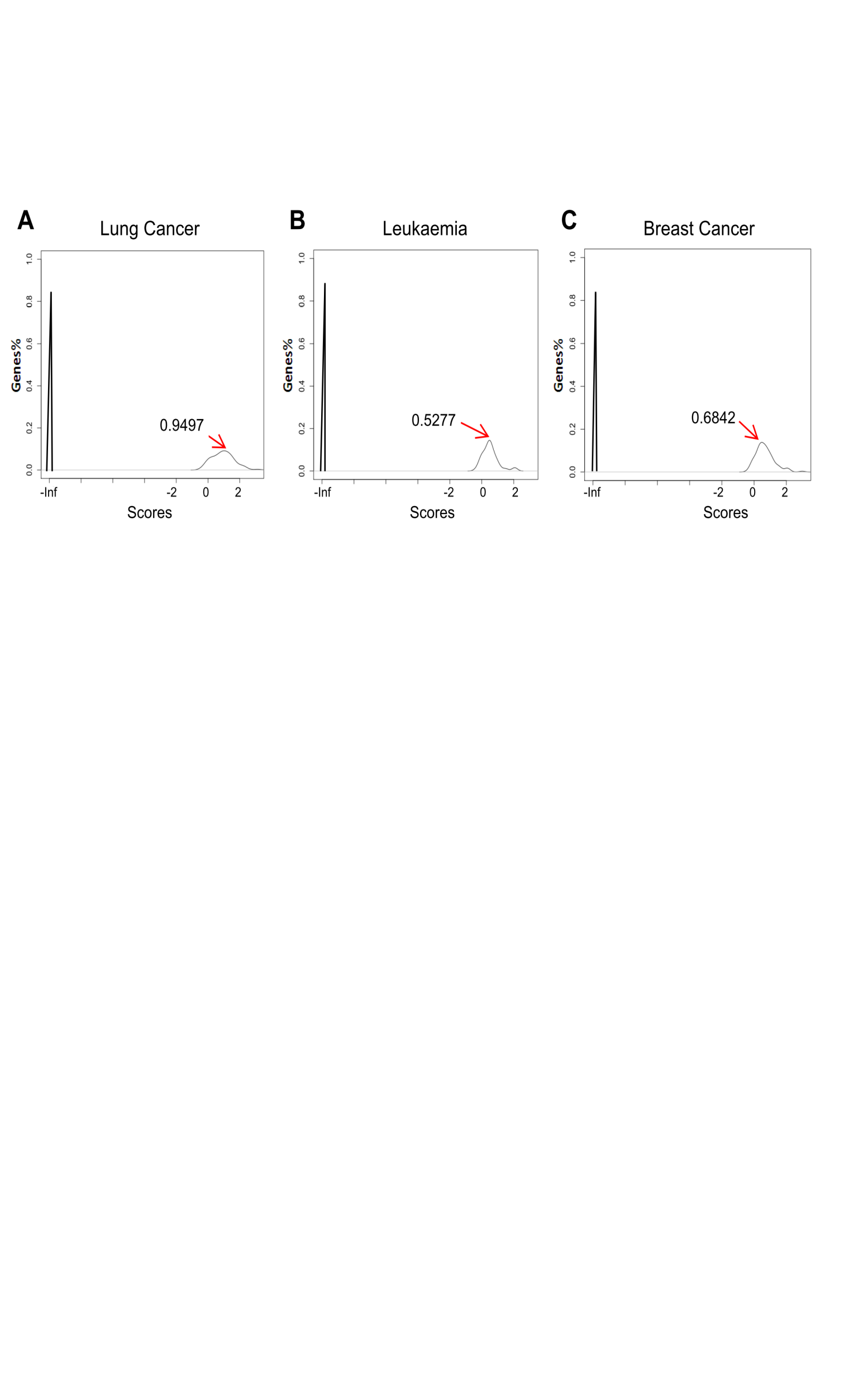
# Additional Figures

## SF1 Co-expression networks constructed by the rank-based method follow the power law distribution

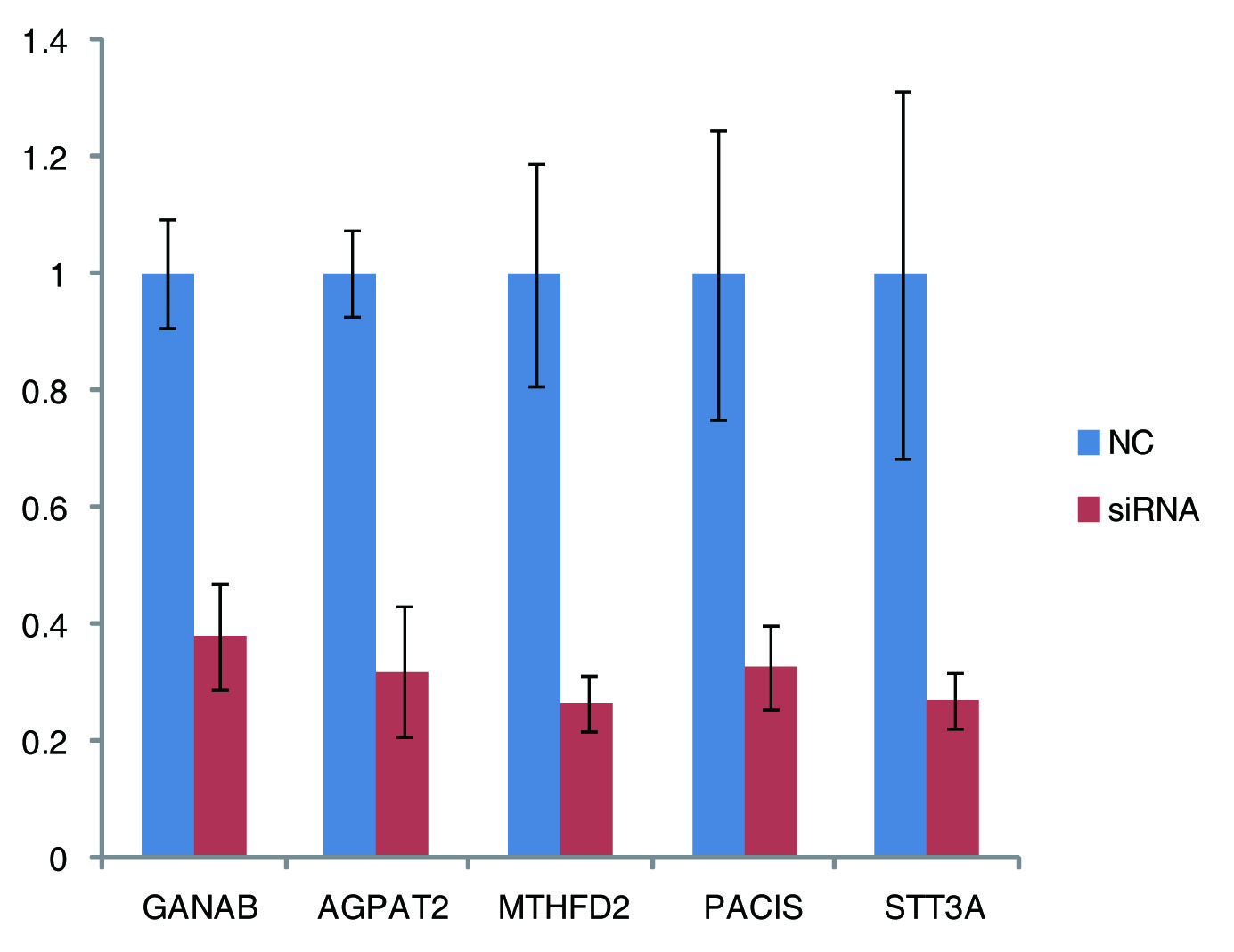
## Figure 1 png.pngSF2 - Genes in cancer-specific modules are enriched in GO terms that associate with cancer development and progression.

## SF3 The distribution of reaction numbers of metabolites and degree centrality in enzyme network.

## SF4 Importance scores in each type of cancer.



## SF5 siRNA efficiency test results.



# Additional Tables

## ST1.pngSTable 1 - The detailed information of datasets used in our analysis

## STable 2 - Cancer Hall Markers and our prediction results

