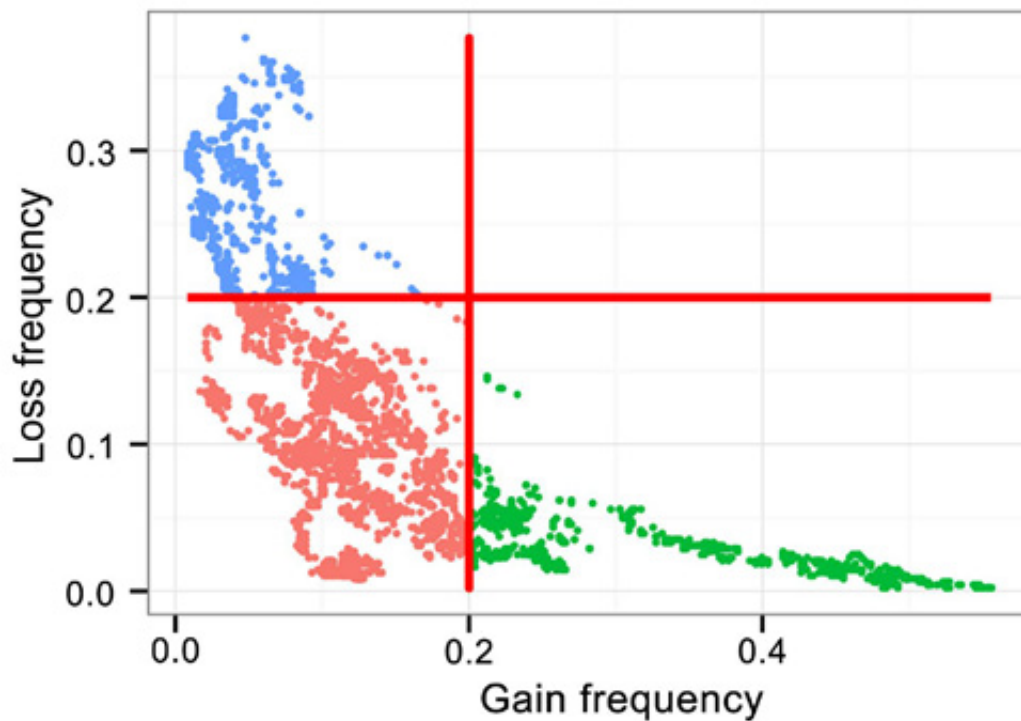
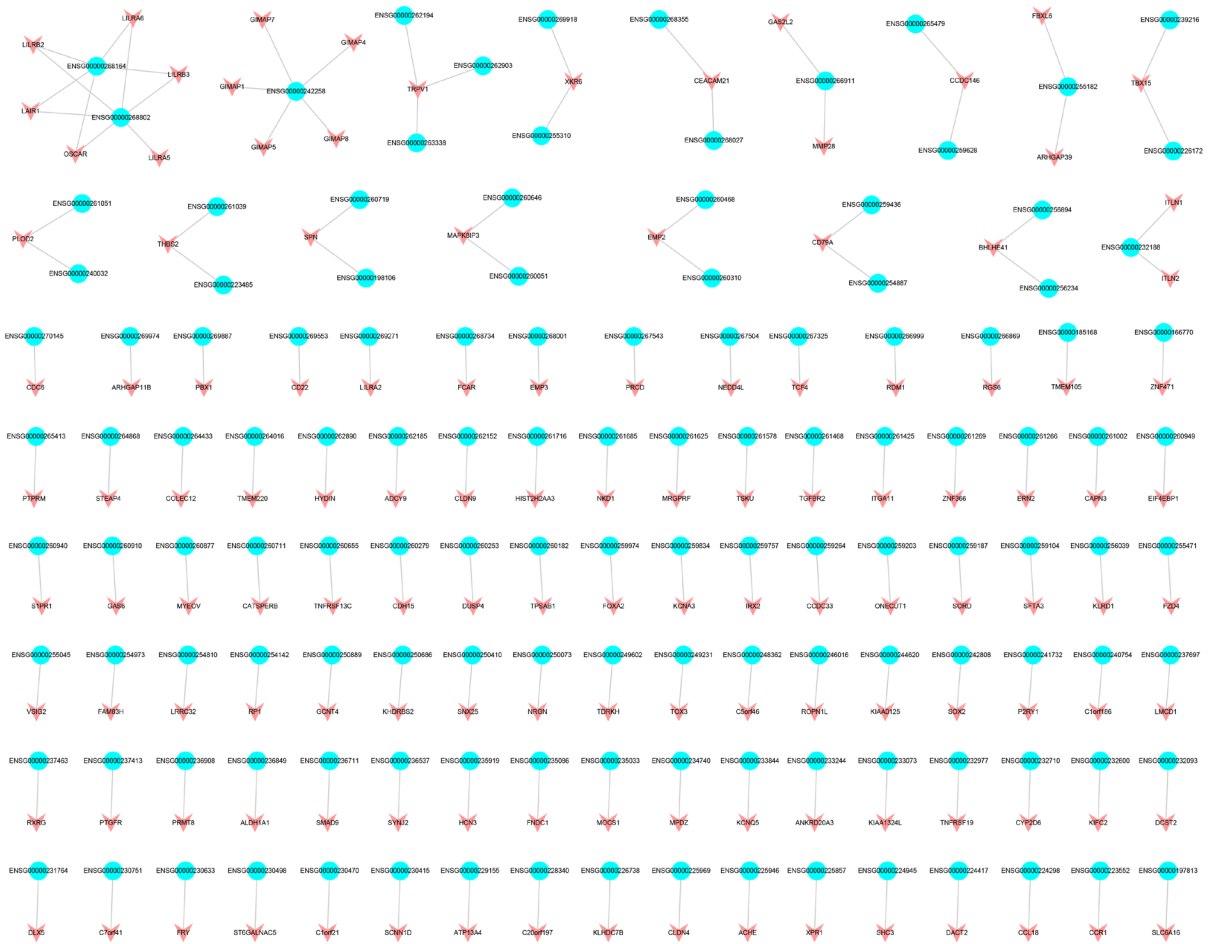


Integrated analysis of dosage effect lncRNAs in lung adenocarcinoma based on comprehensive network

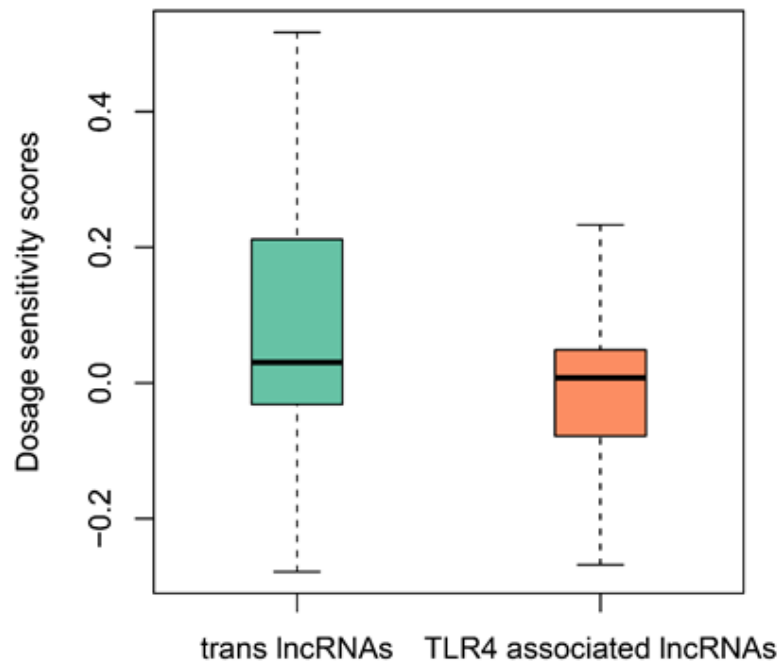
SUPPLEMENTARY MATERIALS



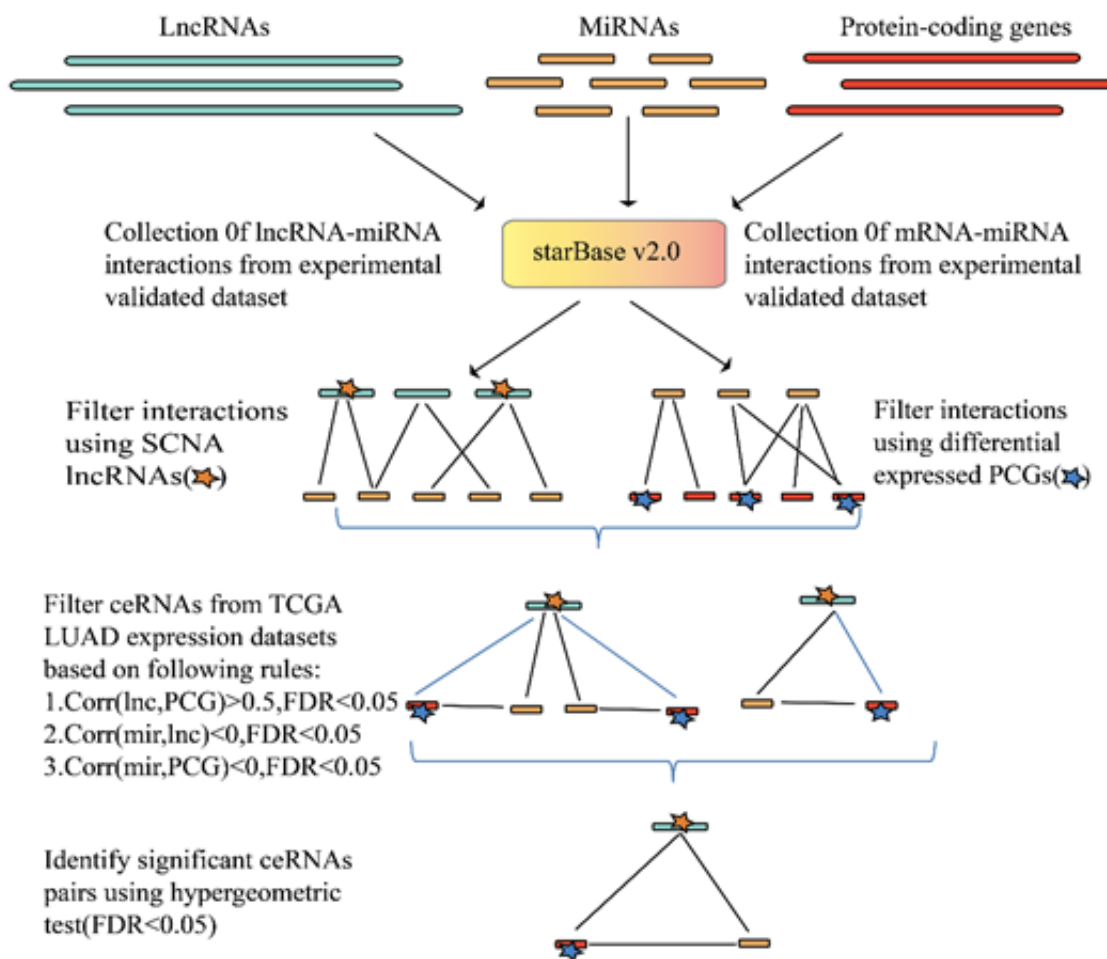
Supplementary Figure S1: The gain/loss frequency of lncRNAs across tumor samples.



Supplementary Figure S2: The cis-acting network of SCNA lncRNAs.



Supplementary Figure S3: The DSSs of SCNA lncRNAs involved in trans-acting network.



Supplementary Figure S4: Pipeline for transcriptome-wide identification of significant LUAD-related ceRNA pairs.

For Supplementary Tables see in Supplementary Files

Supplementary Data S1: 179 amplified and 881 deleted lncRNAs.

Supplementary Data S2: 3,499 SCNA lncRNAs.

Supplementary Data S3: 66 hypermethylation and 2 hypomethylation lncRNAs.

Supplementary Data S4: The PCC between methylated value and expressed value of 17 SCNA lncRNAs.

Supplementary Data S5: List of the potential small molecule drugs.