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Supplemental Information

Upregulated circRNA ARHGAP10 Predicts an Unfavorable Prognosis in NSCLC through Regulation of the miR-150-5p/GLUT-1 Axis Mingming Jin, Chunzi Shi, Chen Yang, Jianjun Liu, and Gang Huang

Supplementary materials 1



Sig pathway of DE gene

Figure legend: Metabolic pathway analysis.

Supplementary materials 2



Figure legend: Agarose gel electrophoresis detection show the target segment of circARHGAP10 or linear RNA treatment with or without RNase R.