

***In silico* prediction of housekeeping long intergenic non-coding RNAs reveals *HKlincR1* as an essential player in lung cancer cell survival**

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Supplementary Tables

Supplementary Table S1

List of HK-lincRNA and TS-lincRNA identified from the HBM RNA-Seq dataset.

Supplementary Table S2

List of 'core essential' HK-lincRNAs.

Supplementary Table S3

Differentially expressed protein-coding genes on knockdown of HKlincR1.