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## **Supplemental Information**

## miR-409 Inhibits Human Non-Small-Cell Lung

### **Cancer Progression by Directly Targeting SPIN1**

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# Figure S1



Figure S1. Screening for potential SPIN1-targeting miRNAs. Immunoblot analysis of the A549 cells transfected with empty vector, miR-23-3p, miR-409, miR-431-5p, or miR-489. Histograms on the right of the immunoblot graphs show corresponding expression levels of miRNAs by qRT-PCR. \*p < 0.05 and \*\*p < 0.01

## Figure S2



Figure S2. miR-409 regulates SPIN1 expression in a posttranscriptional way. Analysis of A549, H460 cells transfected with scramble or miR-409 mimics. The corresponding miRNA expression was determined using qRT-PCR. Each bar represents the mean of at least three independent experiments performed in triplicate  $\pm$  standard deviation. \*p < 0.05 and \*\*p < 0.01.

#### Figure S3



Figure S3. miR-409 regulates the activation of PI3K/AKT pathway. (A) Immunofluorescent staining were conducted in A549 cells transfected with miR-409. (B) CREB1 expression levels were assessed by qRT-qPCR. (C) BCL2 expression levels were assessed by qRT-qPCR. (D) Immunofluorescent staining were conducted in A549 cells transfected with anti-miR-409. (E) CREB1 expression levels were assessed by qRT-qPCR. (F) BCL2 expression levels were assessed by qRT-qPCR. (G) Immunofluorescent staining were conducted in A549 cells transfected with anti-miR-409. (E) CREB1 expression levels were assessed by qRT-qPCR. (F) BCL2 expression levels were ass



Figure S4. SPIN1 is upregulated in NSCLC patients. SPIN1 expression was analyzed in human lung cancer tissues and adjacent tissues from 3 representative donors by Western blotting.

Characteristics	n (%)	
Age		
≥ 60	47 (55.3%)	
< 60	38 (44.7%)	
Gender		
Male	62 (72.9%)	
Female	23 (27.1%)	
Tumor size (cm)		
> 3	34 (40.0%)	
≤ 3	51 (60.0%)	
TNM Stage		
Ι	43 (50.6%)	
ΙΙ	31 (36.5%)	
III	11 (12.9%)	
Pleural invasion		
NO	68 (80.0%)	
Yes	17 (20.0%)	
Metastasis		
NO	51 (60.0%)	
Yes	34 (40.0%)	

Supplementary Table 1 Clinicopathological characteristics of NSCLC patients

Supplementary Table 2 Primer sequences of Oligonucleotides

Name	Forward (5'→3')	Reverse (5'→3')
Primer sequences for real-time quantitative RT-PCR		
SPIN1	GTTCTGGACCAGGTGCCTGTAAA	CTGTCTCAAACATATGTTCCACTG
β-actin	ATCACCATTGGCAATGAGCG	TTGAAGGTAGTTTCGTGGAT
Primers sequences for PCR		
SPIN1 3'UTR	ATGTCATCACAAACTCTGCCAAA	TTTACAGTCAGGAGTTGCACATAAA
SPIN1 3'UTR Mut	CTTGTTCCAGGAATTACCACGGA	TCCGTGGTAATTCCTGGAACAAG
SPIN1	ATGAAGACCCCATTCGGAAAGAC	CTAGGATGTTTTCACCAAATCGTA