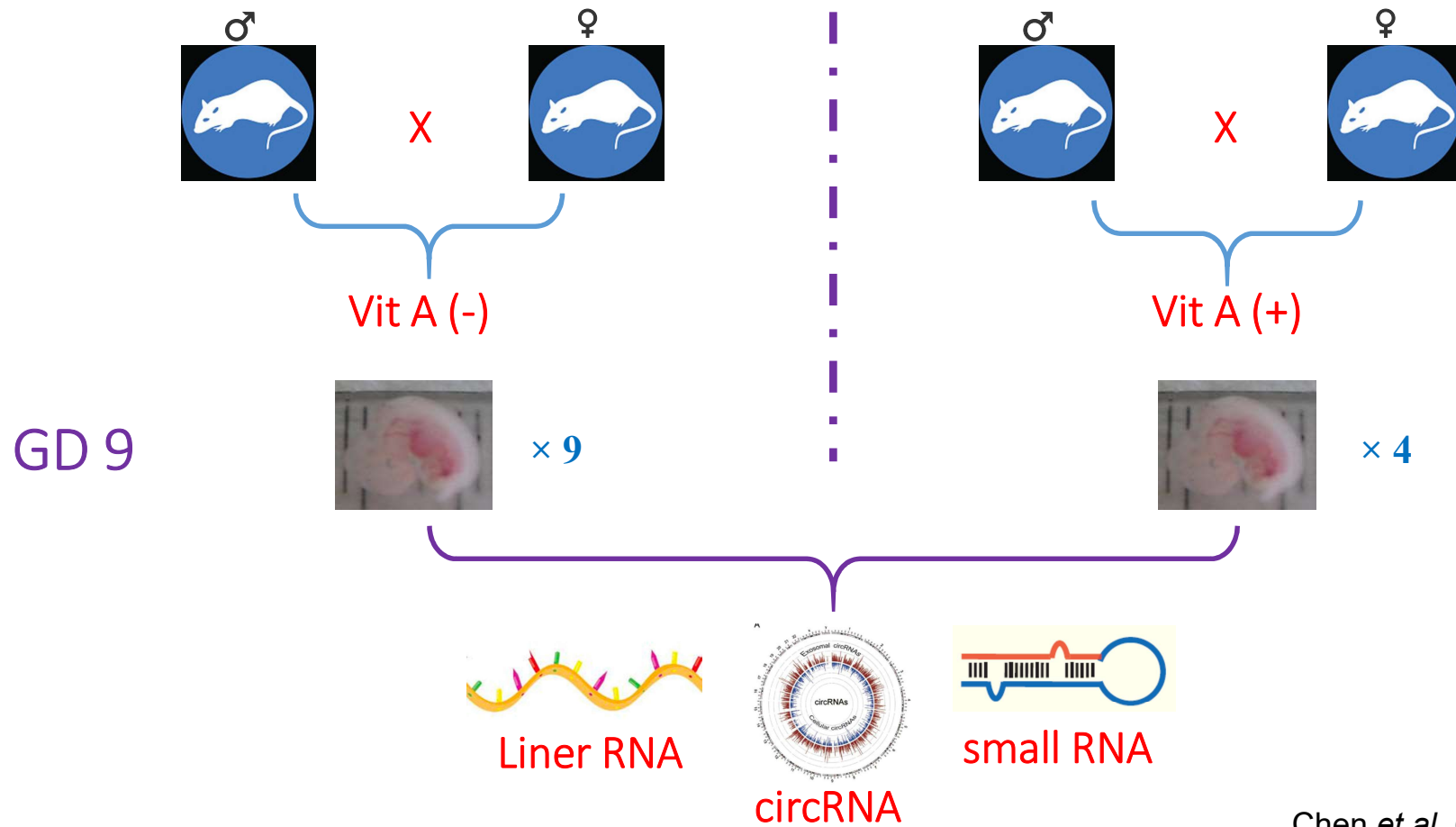


Functions and related mechanisms of lncRNA SULT1C2A in congenital scoliosis

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Transcriptome Assembly

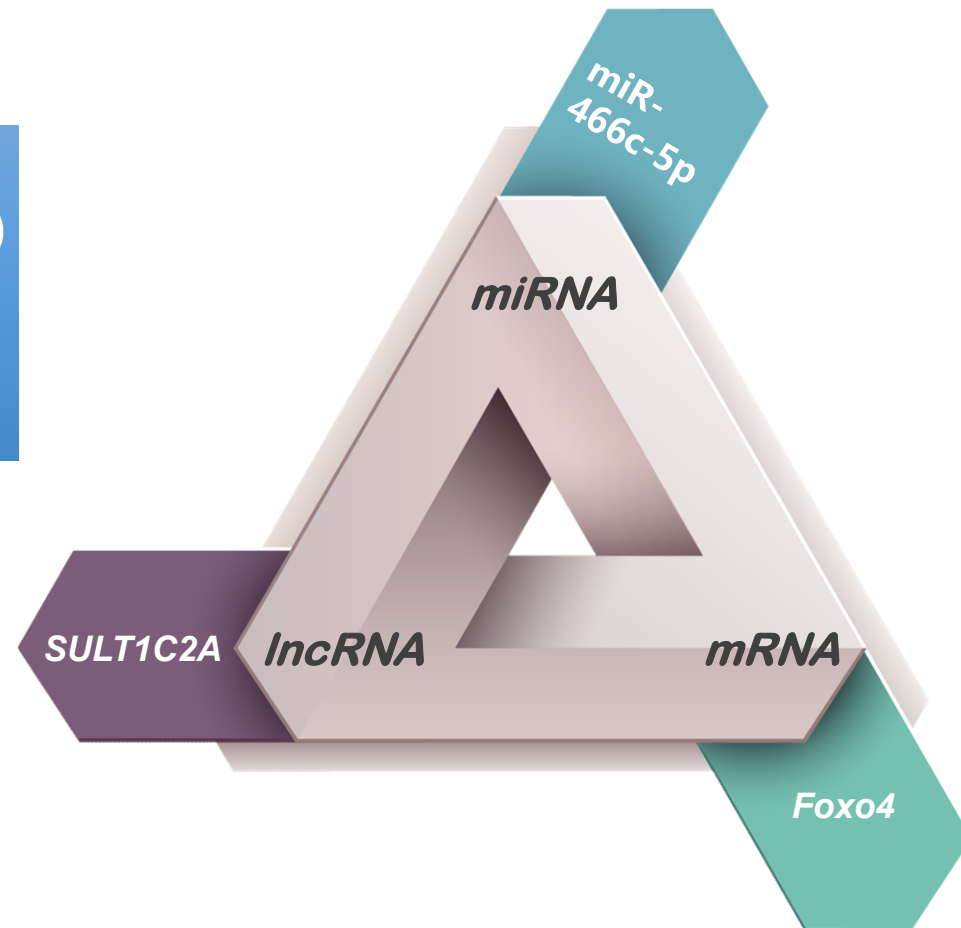


Co-expression network construction

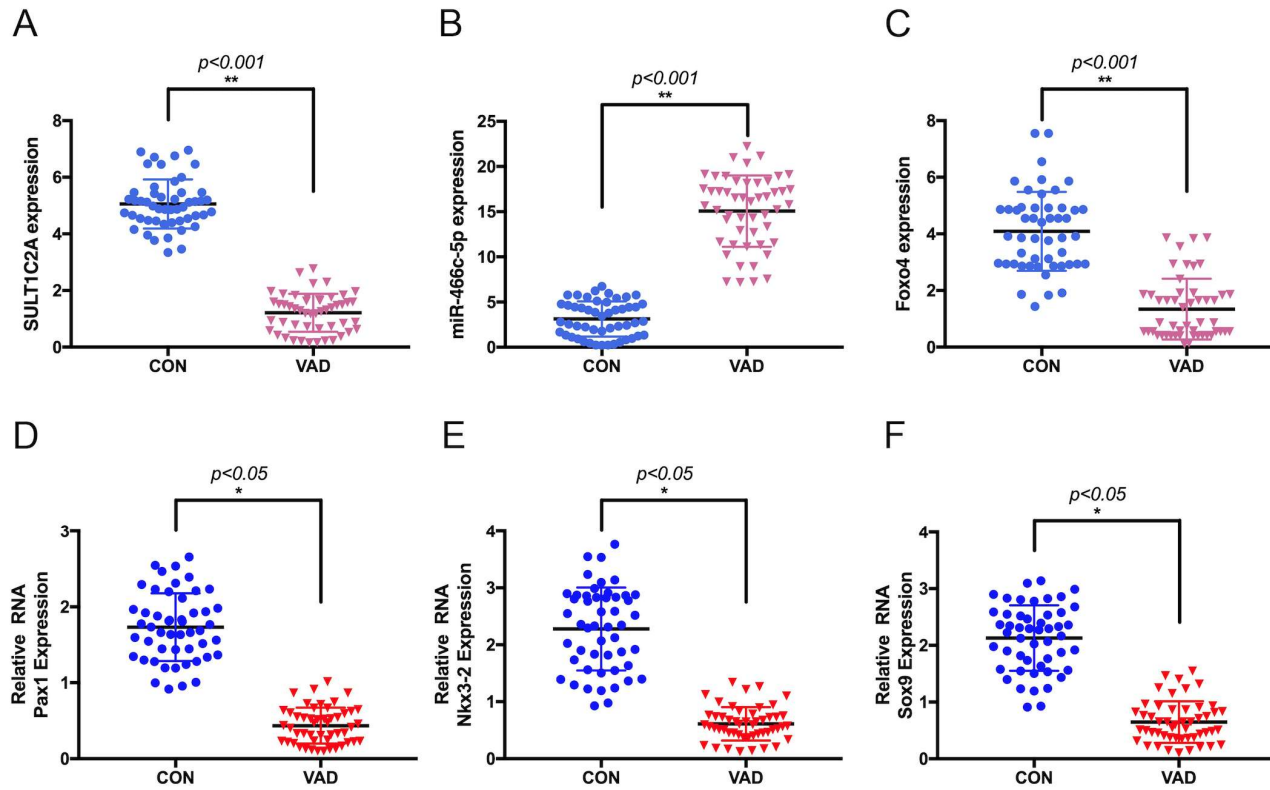
NONRATG027649.1
chromosome 9 (4621424 -4624425 [+])

- intron 4
- Length: 1,828 bp
- Gene: Sult1c2a

Weighted gene co-expression
network analysis (WGCNA)



Expand Sample Verification

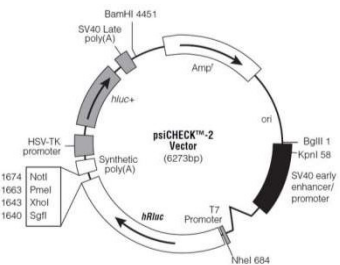
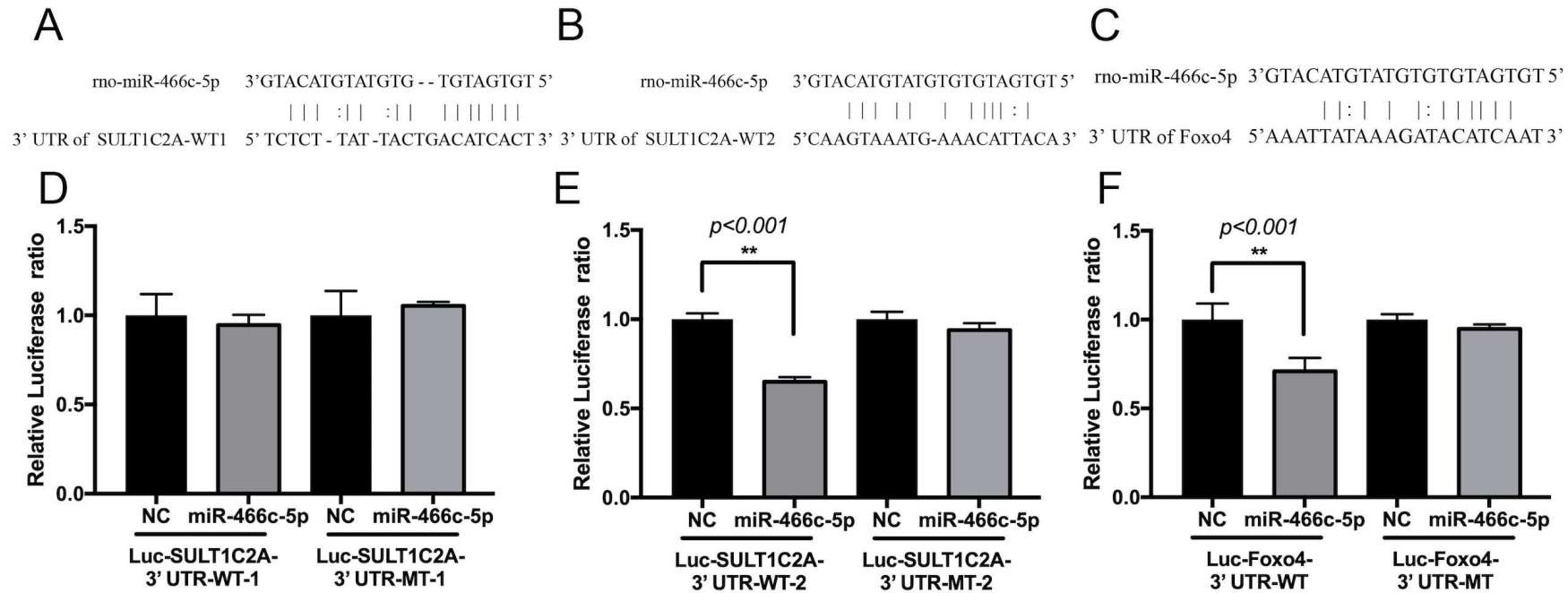


GD 9, n=50

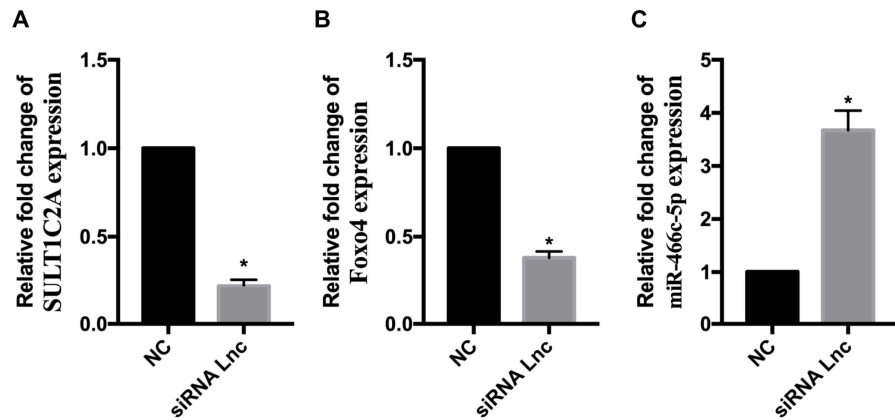
* $P < 0.05$ ** $P < 0.01$. VAD vs Control

Gene	Primer Sequence (5' - 3')
SULT1C2A-F	TGCCCCAGAATGAGAGGTTTGATGA
SULT1C2A-R	CGGTGTCACAGTCCTTGGCATTTAC
FOXO4-F	GGCGGCAAGGGTGGCAAGG
FOXO4-R	CCGGCCTCATTGGGGACAGC
Pax1-F	CAATGCCATCCGCCTACGAAT
Pax1-R	AGAGACCCGCAGTTGCCTA
Nkx3-2-F	AGCGCCGCTTTAACCATCA
Nkx3-2-R	GCGTTTGGTCTTGTAGCGAC
Sox9-F	AGTACCCGCATCTGCACAAC
Sox9-R	ACGAAGGGTCTCTTCTCGCT
GAPDH-F	TGGGGTGATGCTGGTGCTGAGTAT
GAPDH-R	AGCGGAAGGGGCGGAGATGAT
rno-miR-466c-5p-F	TGTGATGTGTGTATGTAC
U6-F	CTCGCTTCGGCAGCACA

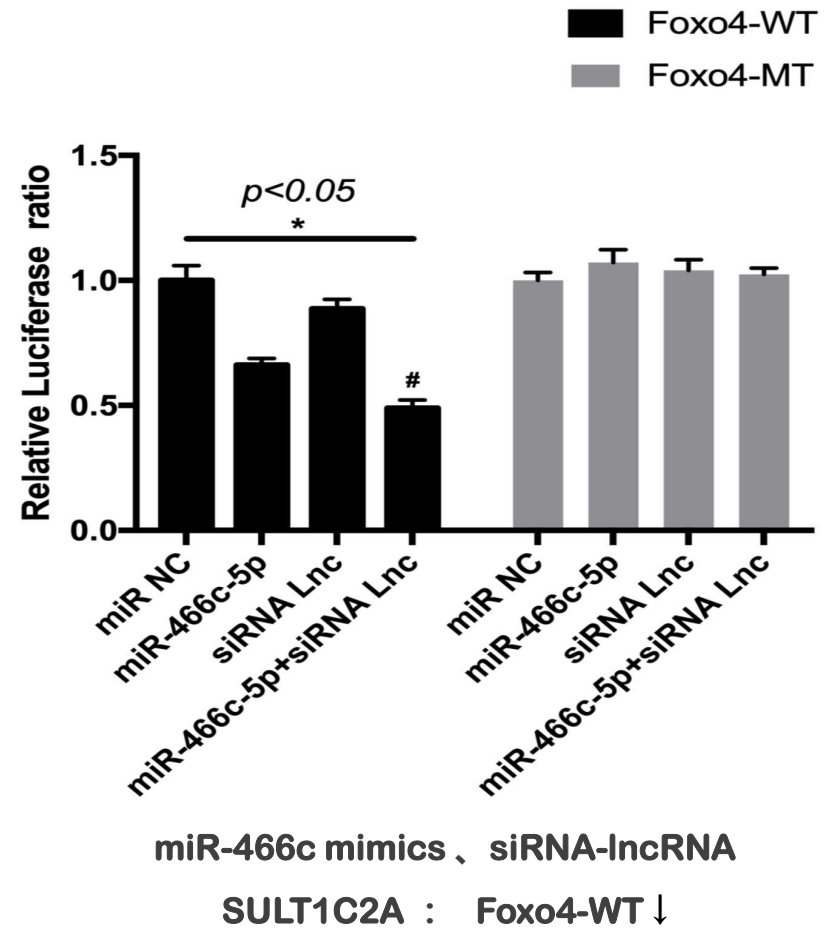
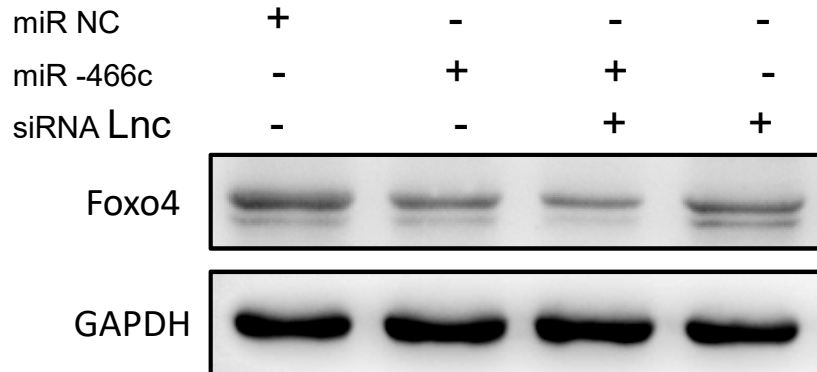
Luciferase Reporter Assays



siRNA Experiments

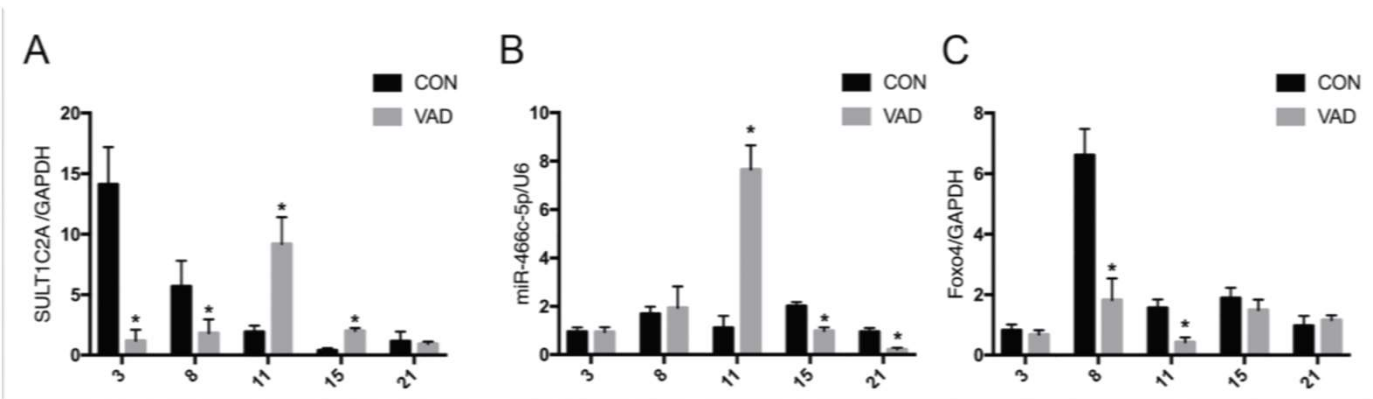


siRNA Lnc : SULT1C2A and Foxo4 ↓,
miR-466c-5p ↑



SULT1C2A-miR-466c-*Foxo4* axis

qRT-PCR



Foxo4



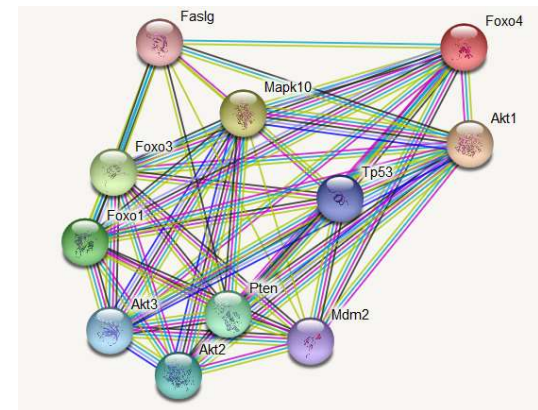
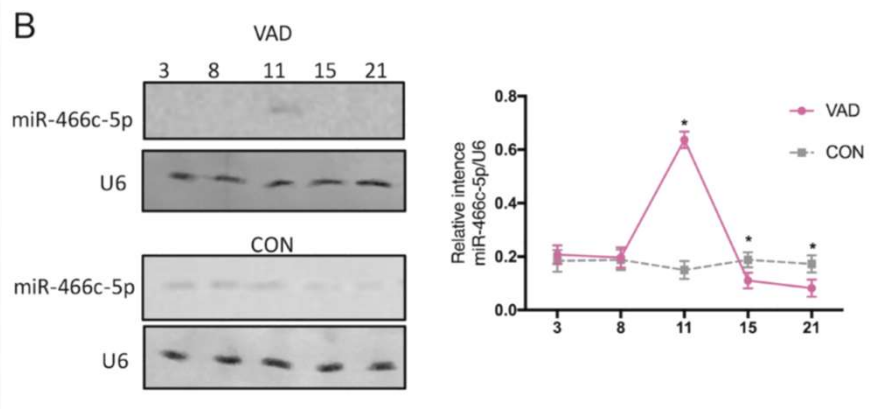
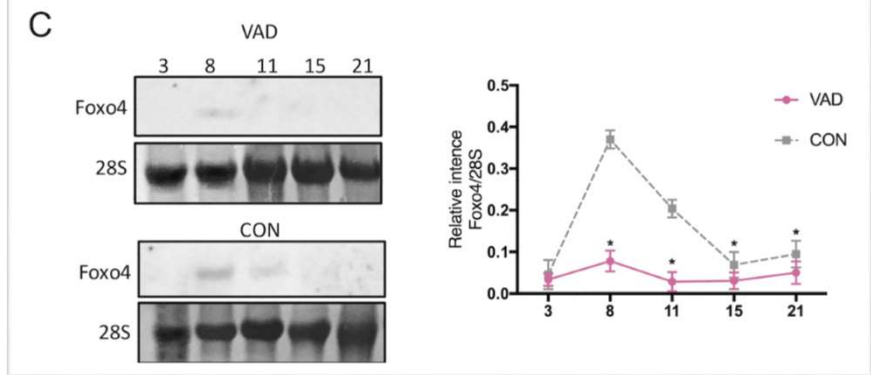
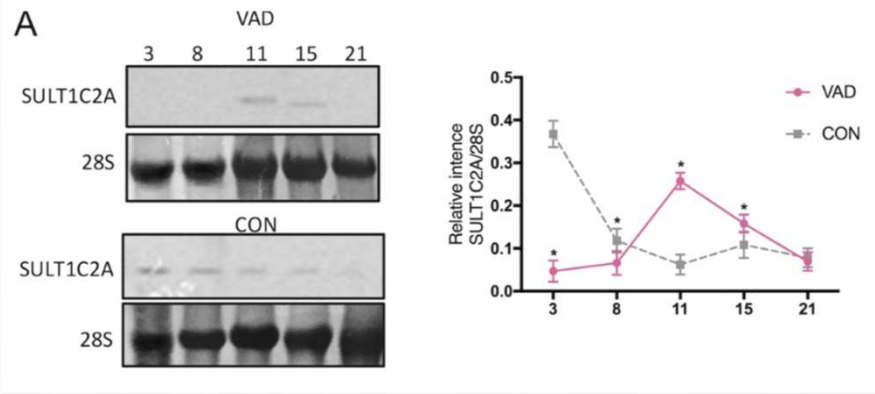
GD 11.5
EMAGE:1315

GD 15.5
EMAGE:13835

- SULT1C2A : GD 3 , 8 ↓, GD **11** , 15 ↑;
- miR-466c-5p : GD **11** ↑ , GD 15 , 21 ↓ ;
- Foxo4 : GD 8 , **11** ↓ ;

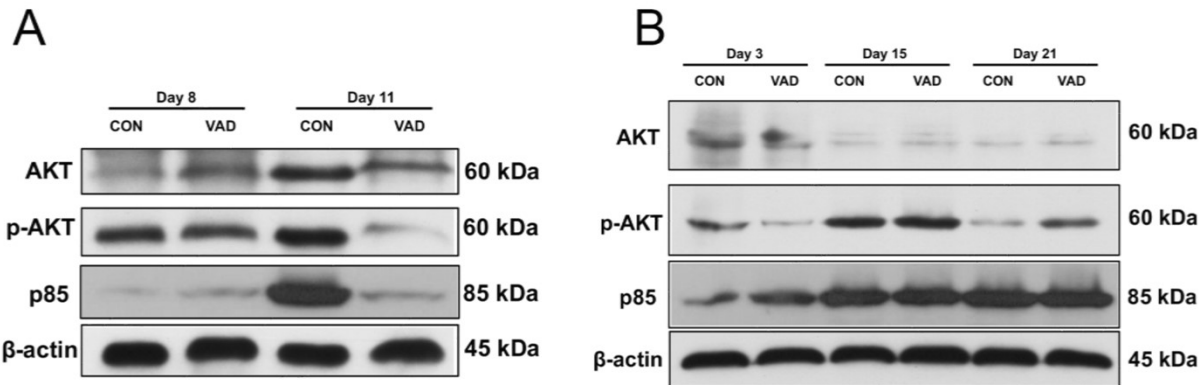
SULT1C2A-miR-466c-*Foxo4* axis

Northern Blot

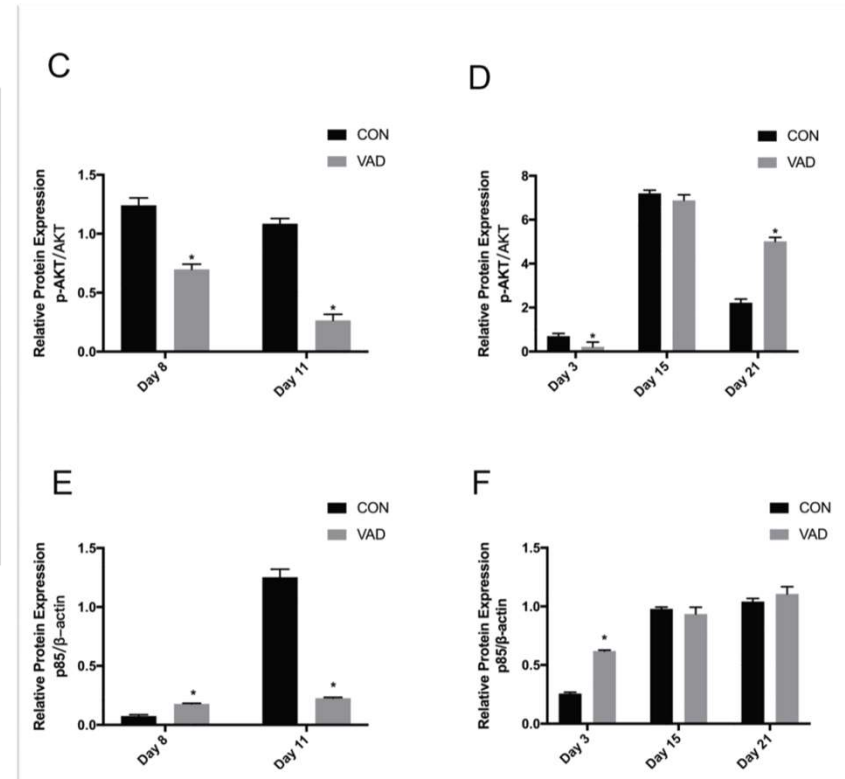


PI3K-AKT-Foxo4 pathway

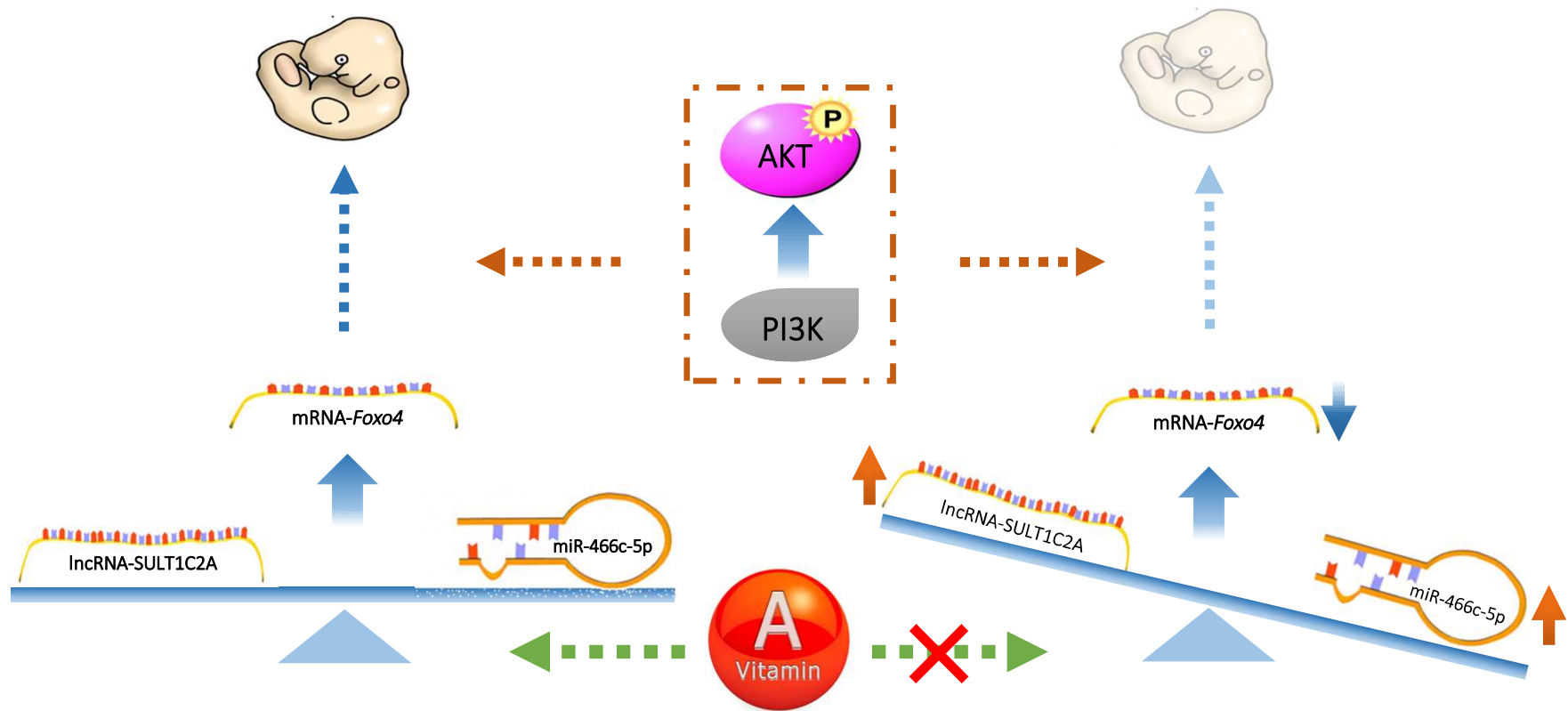
Western Blot



- p-AKT : GD 3 , 8, **11** ↓ , GD 21 ↑ ;
- p85 : GD **11** ↓ , GD 3 , 8 ↑ ;



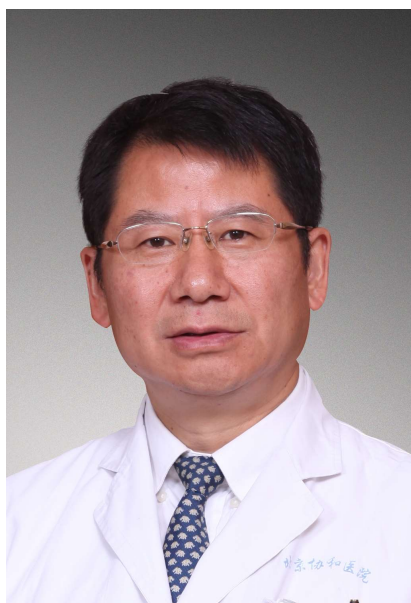
Schematic Diagram of SULT1C2A-rno-miR-466c-5p-Foxo4 axis



GD 9 → GD 11

Conclusions

- VAD initially reduces the expression of *Foxo4* and subsequently up-regulates miR-466c-5p ;
- To maintain the relative balance of embryonic development, lncRNA SULT1C2A is up-regulated on GD 11 and functions as a sponge to antagonize miRNAs ;
- The dynamic changes in the expression of the SULT1C2A-rno-miR-466c-5p-*Foxo4* axis may be initiated on GD 9 and were most obvious on GD 11 during the early-mid stage of somitogenesis ;
- Dynamic profile of AKT phosphorylation, an important signaling pathway for somitogenesis, was found in the VAD group, and this novel finding warrants further research ;



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Thanks