

LncRNA Slfn5os Regulates the Survival and Testosterone Production in Tm3 Leydig Cell Lines

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Abstract

Long non-coding RNAs (lncRNAs) have been reported to regulate the spermatogenesis. In this study, we aim to characterize the expression pattern and roles of lncRNA schlafen 5, opposite strand (Slfn5os) in the testis of adult

Polymerase Chain Reaction (RT-PCR). The localization of lncRNA Slfn5os and Slfn5 was determined by Fluorescence In situ Hybridization (FISH). TM3 Leydig cell line was used as a cellular model to study the function of Slfn5os. The survival of TM3 cells upon Slfn5os knockdown or overexpression was assessed by Cell Count Kit-8 (CCK-8) viability

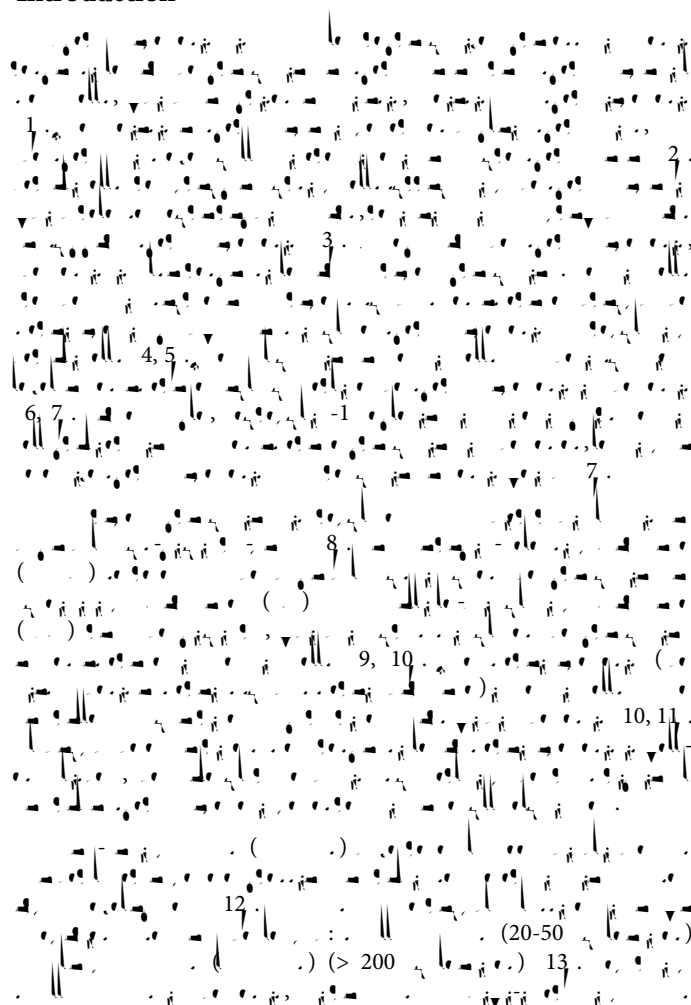
of Slfn5os. The expression level of Slfn5os negatively regulates the mRNA level of Slfn5. In addition, forced expression of Slfn5os impaired the survival and testosterone production in TM3 cells, while Slfn5os silencing showed the opposite

survival and testosterone production in Leydig cells.

Keywords:

lncRNA; Slfn5os; spermatogenesis; testosterone; TM3

Introduction



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Slfn5os (schlafen 5, opposite strand, 10003F1Rik) is a novel lncRNA identified in mouse testis, and it has been found to be regulated by an endocrine disruptor, Diethyl Hexyl Phthalate (DEHP) [20]. It is transcribed from the opposite strand of Slfn5 (Schlafen family member 5) gene. However, its tissue-specific expression and functional role in testis is largely unknown. Elucidating its expression pattern and role in testis can provide insights into the novel mechanism of

Material and Methods

Animals and cell culture

57 /6
12- /12-
3
-2. (.)
1:1 () : 12
1% / 5% () 37+
5% -2
10% () 1% /

Cell transfection

5 () 5 ()
3.1- 3.1- 5
3 6- 1 10⁵
24
2000 ()
3 6- 60%
50 6 8
48

Fluorescence *In situ* Hybridization (FISH)

3 4° 4%
5
21
(1:500)
3 (1:50)
1:100
5
5- :5 -
-3-(30
5: 5- -3-(25

Quantitative RT-PCR and RT-PCR (reverse-transcription PCR)

()

(2)
()
7500
()
95° 2
40 95° 30. , 60° 30. 72° 60. 2
5- 5
42 94+ 45
57° 1 72° 90
1%
1,
5- 5
45

Cell proliferation assays

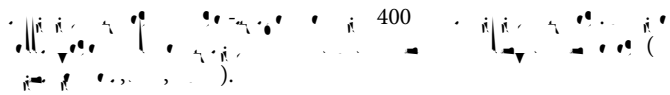
8 (8;
96
5000 48, 10 8
1 37°
10
490
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Apoptosis detection by flow cytometry

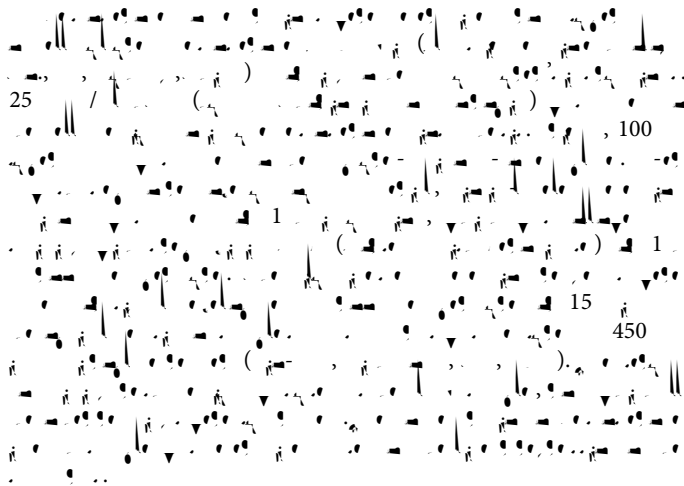
3
(4
1 106 / 5 5
30

Table 1: Primer sequences.

Gene	Primer Sequence
Slnf5 (RT-PCR)	
Slnf5os (RT-PCR)	
Slnf5 (qPCR)	
Star (qPCR)	
Cyp11a (qPCR)	
Lhr (qPCR)	
Hsd3b1 (qPCR)	



Testosterone measurement by ELISA (Enzyme-Linked Immuno Sorbent Assay)



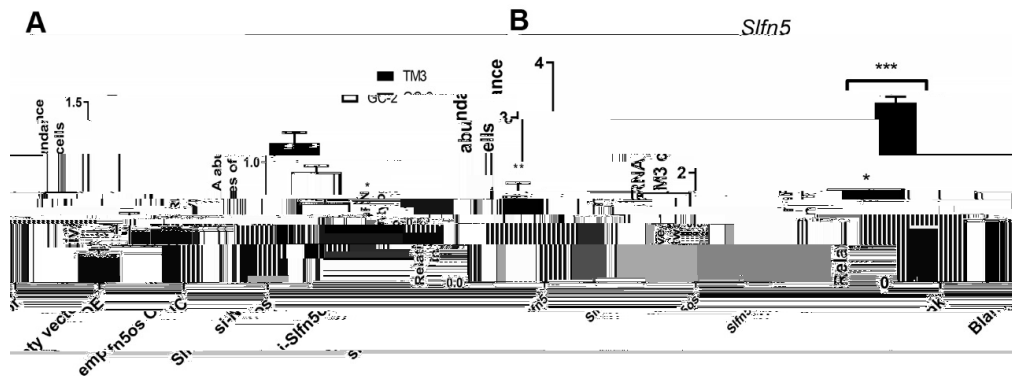
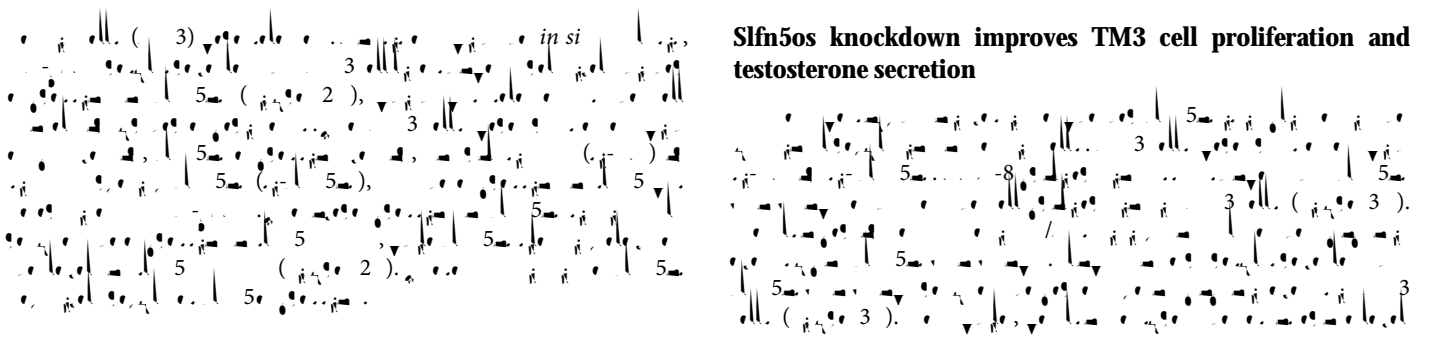


Figure 2: The impact of Slfn5os on Slfn5 expression. (A) The expression levels of Slfn5os on Slfn5 were examined by RT-qPCR in TM3 cells (mouse Leydig cell line) and GC-2 cell line (mouse spermatocytes). (B) mRNA expression of Slfn5 in TM3 cells after the transfection of Slfn5os empty vector, Slfn5os expression plasmid, si-NC or si-

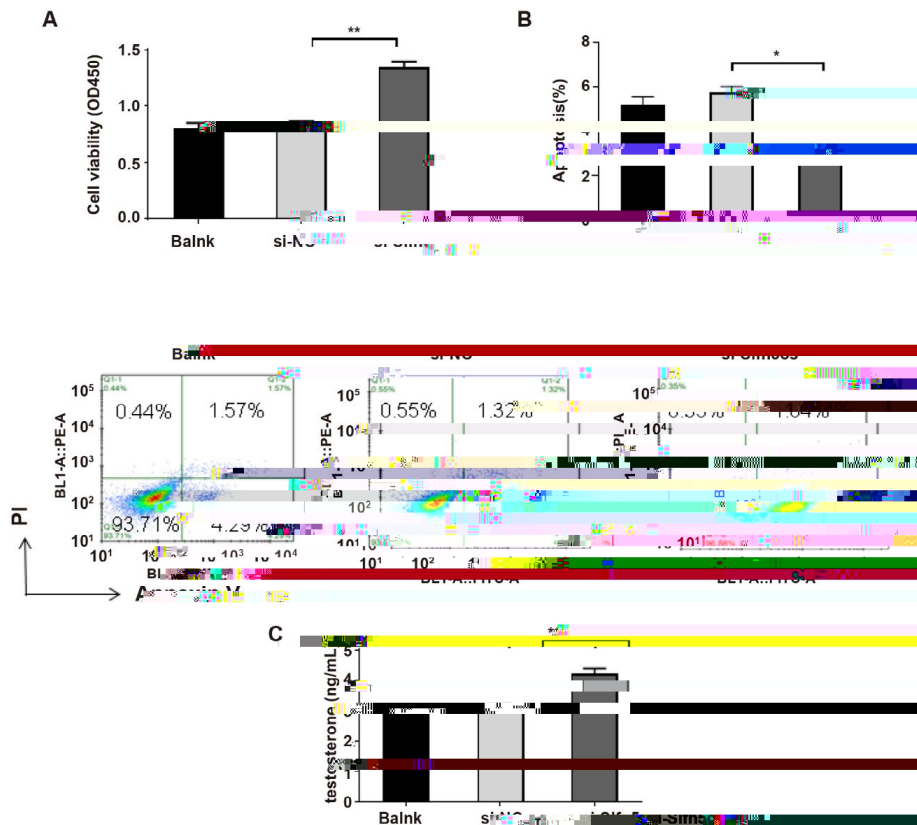


Figure 3: Slfn5os knockdown promotes TM3 cell proliferation, inhibits cell apoptosis and increases testosterone secretion. (A) CCK-8 proliferation assay in

5 (3).

Slfn5os overexpression impairs cell proliferation and testosterone secretion

5 3 5 5 3 (4), 5 (4), 5 (4), 5 (4), 3 5

Slfn5os modulates the expression of steroidogenic genes and enzymes in TM3 Leydig cells

5 3 5 3

11 1, 17 1, 3 1, 17 3 11 1, 17 1, 3 1, 17 3 5 (5), 11 1, 3 1 3-7 5 (450) 11 1 3 5 11 1 5 11 1 5 11 1 5 11 1

Consent for Publication

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Competing Interest

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Author's Contributions

Acknowledgement

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