

# Knockdown of TOR causing ovarian diapause in a genetically stable brachypterous strain of *Nila a a al gen*

Fangzhou Liu<sup>1</sup>  | Kaiyin Li<sup>1</sup> | Wanlun Cai<sup>1</sup> | Jing Zhao<sup>1</sup> |  
Yulan Zou<sup>2</sup> | Hongxia Hua<sup>1</sup>

<sup>1</sup>H b il R U iiia a  
S aiab P Ma a K  
Lab a ,C Pa  
Sci a T ,  
Ha A aU i i,W a ,C ia  
<sup>2</sup>C Li Sci a T ,  
Ha A aU i i,W a ,C ia

**Correspondence**  
H b H a,C Pa Sci a  
T ,Ha A aU i i,  
W a 430070,C ia.  
E ai: a @ ai. a .  
G a :Na i a Na a Sci F a-  
i C ia;G a b :31171846.

## Abstract

B a (BPH), *Nilaparvata lugens* (S ) (H i a: D - aia ), i a a i BPH i a ia i a a a a a i a a i a a .T a a i a i a a a a i a i i

I i i a i i , i i a i a i BPH.A i i a i a a  
 i MF, i a i i a i i a i BF(J , 1963; Kii , 1956).  
 I abia i a ab , BPH i BF i a i a i a i (S b ,  
 Mi i a, Ya a Ma a i&Ma a, 2002); abia i i , BPH i MF  
 i abii a ii a i a i (Kii , 1956).  
 I a i i i , MF a a a i i i i a BF a (Kii , 1965).  
 T i i i a b i a a i a a , a a a i a  
 i i a i (C , C , Ya , & Yi , 1979). BPH i i i b Insulin Receptor 1 a Insulin  
 Receptor 2

**TABLE 1** List of primers used in this study

GenBank accession number	Name of primer	Primer sequences	Length of PCR product
JQ793898.1	N1TOR-F	GATCGGGCATGAGGGAGGGAGACA	6,768 b
	N1TOR-R	CGACGACGGTACACTGCGTTGG	
	1N1TOR-F	TAATACGACTCACTATAAGGACCAG TGAAATGCTCGTAAACA	550 b
	1N1TOR-R	TAATACGACTCACTATAAGGCCAGGTGCAGG TAATCGTCCAG	
	2N1TOR-F	TAATACGACTCACTATAAGGTTGACGG TCACTCACTACTGCA	492 b
	2N1TOR-R	TAATACGACTCACTATAAGGCTCTTGTT TCGTCCCCATACC	
U76561	GFP-F	TAATACGACTCACTATAAGGTAAA CGGCCACAAGTCAG	400 b
	GFP-R	TAATACGACTCACTATAAGGTCGGC CATGATATAGACGTT	
EU179846.1	PCR-N <i>I</i> Actin1-F	CCAACCGTGAGAAGATGACC	256 b
	PCR-N <i>I</i> Actin1-R	GATGTCACCGCACGATTTCAC	
JQ793898.1	PCR-N1TOR-F	AACGCCATGGAGGTGACAGG	143 b
	PCR-N1TOR-R	ATGAGGCGCCAGTTGAGCAG	
KY827832	PCR-NIE74B-F	AACAACATAATAGGCACAGTC	175 b
	PCR-NIE74B-R	GGAATGGCGAAGAAGTATC	
FJ263049.1	PCR-N <i>E</i> cR-F	GCCAGAAAGTACGACGTGAA	234 b
	PCR-N <i>E</i> cR-R	TTGGATCTTCTCCACCTTCC	
19JF345255	PCR-N <i>F</i> oxA-F	GCGGAGGTTATGTTGTTGTA	193 b
	PCR-N <i>F</i> oxA-R	CTGAGCCTTGTAGCATGTTGAA	
AB353856.1	PCR-N <i>V</i> g-F	TCTTCATCATCCTCCTCCTCTTC	173 b
	PCR-N <i>V</i> g-R	TCCTGGTTGTTGTCATTGTCATT	

### 2.3 | Synthesis of dsRNA

T b - a RNA ( RNA ) i ■ N1TOR. T RNA ( 1N1TOR, 550  
 b ) a b 2,764 a 3,313 ■ i , a ■ RNA ( 2N1TOR, 492 b ) ■ a b  
 1,736 a 2,227 ■ i .T7 a a a a 5'- - ■ i ■ i .T  
 i a i ■ DNA N1TOR. T i PCR ■ ■ a ii i T7 a a a  
 b 5'- a a RNA i .QIA ■ TM PCR i ■ a i i ( C N : 28104, Q a -  
 , D , G a ) i PCR ■ . RNA i i i aT7 R b MAXE  
 RNA i S ( C N : P1700, P a, Ma i ) i a a a ■ ' i ■ i .GFP ( G Ba a ■ -  
 i .U76561) RNA ( GFP ) a a a i ■ .T ■ i i i i RNA a  
 1N1TOR-F/R a 2N1TOR-F/R (Tab 1).

### 2.4 | RNAi using microinjection

T i - i a ( 6 12 ) BPH ■ ■ RNA i ■ i i ■ i ■ a ■ b b Li  
 a. (2015) i a Na i 2000 i ■ (WPI, Sa a a). A i a 300 RNA ( 1N1TOR, 2N1TOR,

## 2.5 | Developmental duration and ovary observation

Ea BPH a i RNA a b a 6 i i a a i i i i N  
 a a a i i - a .T BPH a i i i i i - i i  
 (WPI, Sa a a) a 1,3,5,a 8 a a a , i 1× a -b a i (PBS) b a-  
 i i 4% a i 1× PBS 20 i a a .Di a i a a i  
 10 i i 0.2% T i -X 100 (C N :.T9824, Sa a a, Si a) i 1× PBS. A a i , a i -  
 a i a O (SZX16,O ,T ,Ja a ) a P90 P Di i a Ca a  
 (P90,Ni .T ,Ja a ).R i i BPH i a b .

## 2.6 | qRT-PCR

## 2.7 | Sequence comparison and phylogenetic relationships

P      ba      a i a i      ,      i ,  
i b      i b - i i      i M a E      i a G      A a i (MEGA)      a .  
T b      a a a i i 1,000 a .

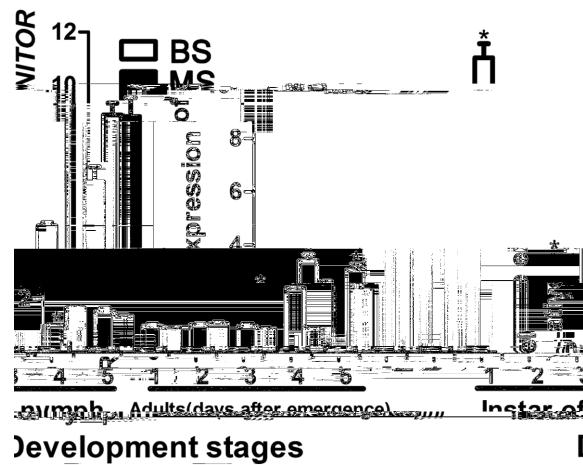
## 2.8 | Data analysis

A a a a a i SPSS i 18.O - a ANOVA a a a i b a ,  
a a - a ANOVA a a a i a RNA.P a a a b  
a i b a i b a a i.

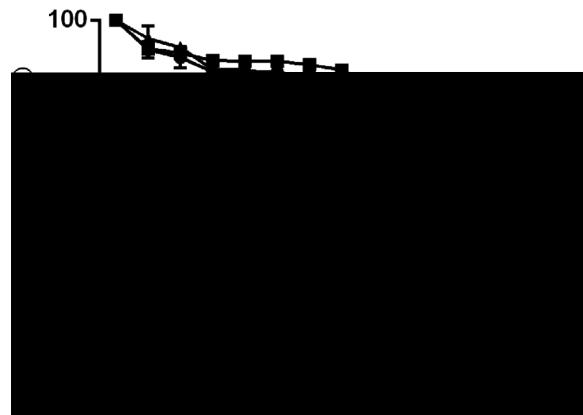
## 3 | RESULTS

### 3.1 | NITOR expression patterns during different developmental stages in BS and MS BPH

T a NITOR a i b a a i BPH; i a i a ai  
i a a .D i a a , NITOR i a i a i BS a  
MS a i a , b i i a i i NITOR i a a a .  
T NITOR i i BS a i i a i MS a a 1,2, a 4 a a



**FIGURE 1** T a i i a NITOR a i a a MS a BS; BPH actin1 a  
 a a a .T i a a i a i a a a BS. T a a  
 i a ba b i a .E ba i a a a .Ba ab i a  
 a i i i i a b BPH MS a BS, a i i - a ANOVA ( $P < 0.05$ )



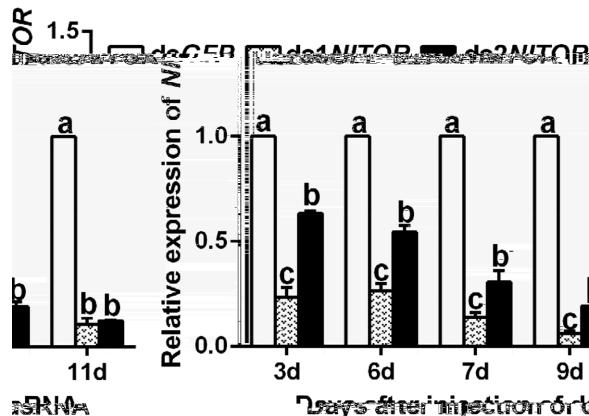
**FIGURE 2** S i a a BS BPH i 1NITOR a 2NITOR i i . GFP i i a a  
 i . Ea i 45 . S i a GFP, 1NITOR, a 2NITOR a a  
 i a - a ANOVA

■ , a NITOR i a i BS a MS 3 a a ■ (F i 1).T NITOR  
i i i i i a a a 5 a ■ b BS a MS (F i 1).

### 3.2 | Effects of knockdown of *NITOR* on the survival of BS BPH

B **a** i NITOR i BS a i i **a** i a i MS a a 1,2,a  
 4 a a **a**, NITOR a i **a** i BS i i a NITOR i **a** i a-  
 a i BPH.

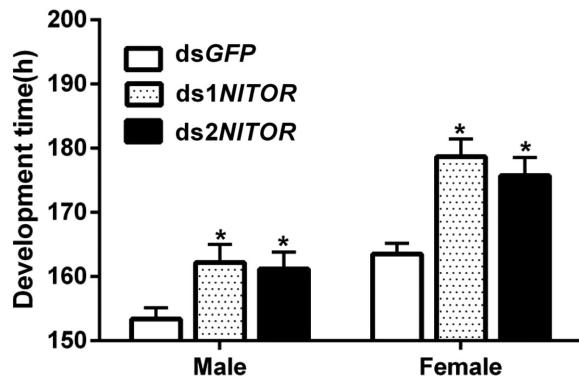
A i **a** i RNA( 1NITOR, 2NITOR, a GFP) i i - i a , i a a  
 a i NITOR a GFP i i 1 7 a a i **a** i (F<sub>2,6</sub> = 0.538; P = 0.609; F i 2). I  
**a** , i a a i **a** a i NITOR a i **a** a a i **a** a i  
 GFP 8 15 a a i **a** i (BS: F<sub>2,6</sub> = 12.568; P = 0.007; F i 2). F 1 7 a a i **a** i , a



BPH i a a , i 8 15 a a i BPH , a BPH i a a . T i  
i BPH a a NITOR a a BPH a b BPH .

### 3.3 | Effects of dsRNA injection on *NfTOR* gene expression in BS

BS, i NITOR a 2NITOR i i - i a i i a NITOR, a i  
 NITOR RNA a a a i i .F 3 11 a a i i NITOR, a i  
 NITOR a i i a b 36.9 94.1% (F) 3 (P <



**FIGURE 4** T 1NITOR a 2NITOR i a i BS BPH  
 i i a a .A ba i a a a a i i i a ab i i i i a i i a a , a i - a ANOVA ( $P < 0.05$ )

### 3.6 | Knockdown of NITOR inhibited ovarian development

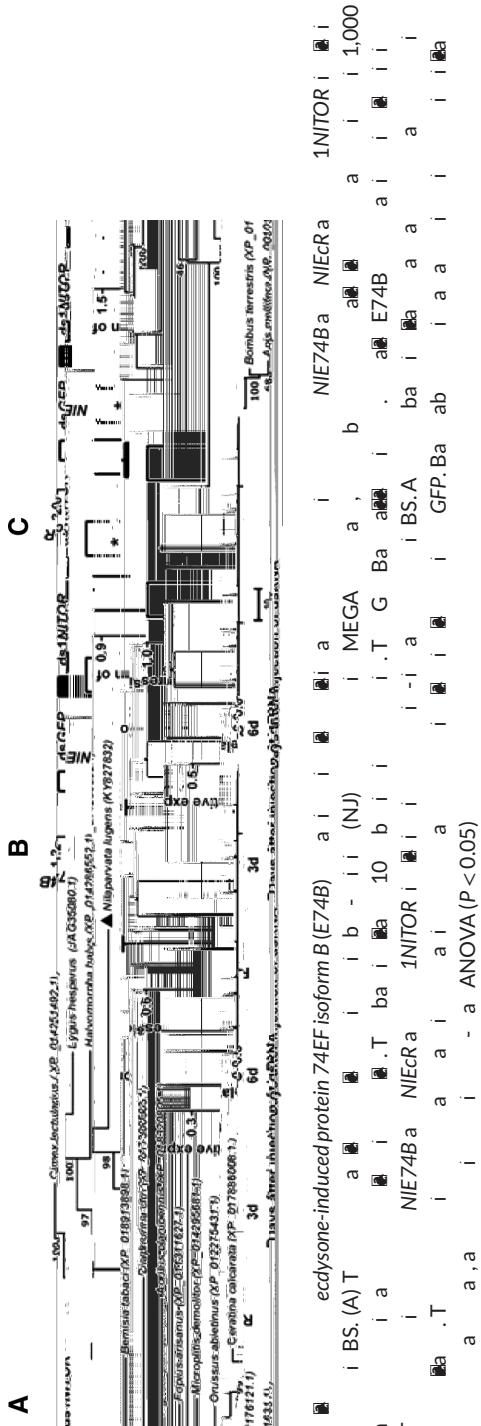
W a a BS i a i 1NITOR a i a a i .  
H , a BS i ■ i GFP a i a a a b a ■ (ab 100  
/ a ).T , i ■ a i a a 1,3,5,a 8 a a ■ a i a  
.T a a ■ a i b i a i ■ a a ■  
a b 1NITOR i ■ , a i a ■ i a ■ b a 3,5,a 8 a A ■ i  
a i ■ i a BPH a ■ b L (2011), a i NITOR- a BS a ai  
i ■ a b a i i .B ■ a , GFP- a a a i -  
a a , a i a b a 3,5,a 8 a (Fi 6). T ■  
a a i BS a i a i NITOR a i ■ i a a a-  
ba C a .(1979).

### 3.7 | Effects of in vivo knockdown of *NITOR* on the expression of *NIFo A* and *NIVg* related to ovarian development

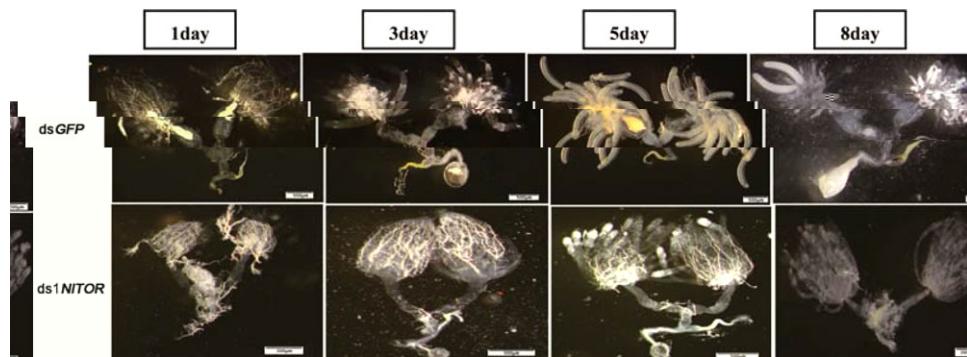
P i i a a FoxA a Vg (Vitellogenin) a i i a a (D a., 2011).  
T i NIFoxA a NIVg i i a b RT-PCR a 1,3,a 5 a a ■ .  
A 1,3,a 5 a a ■ , NIFoxA i i a a i 1NITOR a ■ b 95.9, 89.7,  
a 25.5%, ■ i (Fi 7A) i ■ a i ■ . A 1,3,a 5 a a ■ , NIVg i  
i a a i 1NITOR ■ a i i ■ a b 71.6, 60.8, a 51.2%, ■ i (Fi 7B), ■ a  
■ (P < 0.05, - a ANOVA).

## 4 | DISCUSSION

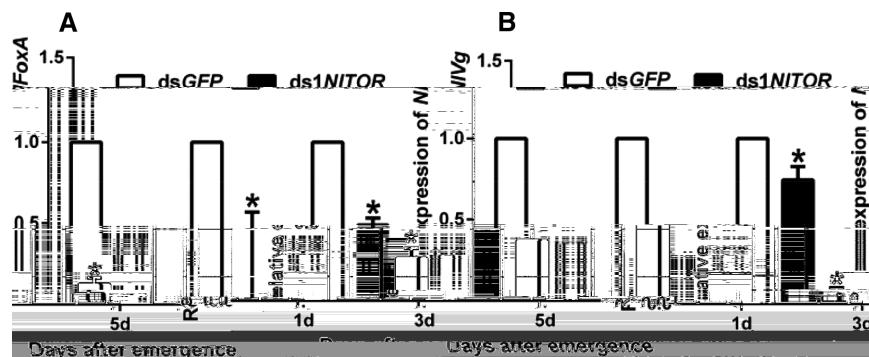
TOR (O a., 2000; Z a., 2000). I (L a., 2016; Ma a., 2009; Pa a., 2006). NITOR a b (L a., 2016; Z a., 2015; Z a., 2017). D i a i , a BPH i a a i i - a , a i , (C a., 1979). W a i a i b NITOR a BPH i a a ai . I , a NITOR i



**FIGURE 5** Phylogenetic trees of ecdysone-induced protein 74EF isoform B (*E74B*). **A**, *E74B* from *Cimex lectularius*, *Liposcelis heparinus*, *Halictus ligatus*, *Nilaparvata lugens*, *Bemisia tabaci*, *Copostoma sinense*, *Popillia japonica*, *Microtia laetabilis*, *Oncrus abietinus*, *Ceratina ciliata*, and *Trichogramma cacoeciae*. **B**, *E74B* from *Cimex lectularius*, *Liposcelis heparinus*, *Halictus ligatus*, *Nilaparvata lugens*, *Bemisia tabaci*, *Copostoma sinense*, *Popillia japonica*, *Microtia laetabilis*, *Oncrus abietinus*, *Ceratina ciliata*, and *Trichogramma cacoeciae*. **C**, Relative expression levels of *E74B* isoforms in different genotypes. Genotypes are grouped by letter: BS, (A) T, (B) T, (C) T, MEGA, (N), T, G, Ba, E74B, BS.A, GFP.Ba, and NIE74Ba. Statistical significance was determined by ANOVA ( $P < 0.05$ ).



**FIGURE 6** E . 1NITOR a . 1NITOR- a a (300 / i ) a a - a a i b i a 1,3,5,a 8 a a . GFP(300 / i ) a a i a a a a 1,3,5,a 8 a a .



**FIGURE 7** T a i i NIFoxA a NIVg a i i RNA. (A) T a i i NIVg  
 NIFoxA a i i 1NITOR i a i i a .(B) T a i i NIVg  
 a i i 1NITOR i a i i a .E ba i a a .T i i a  
 a i a i a i a i i GFP. Ba ab i a a i i i i a  
 b a a a a a i i - a ANOVA ( $P < 0.05$ ). T a a a i  
 a ba bi a a

a BPha , BPH .W i a a NITOR a a  
 i a i a a i i i i NITOR i a a i  
 b i .  
 P i a a NITOR a a a a a JH a i -  
 a a (NIJHAMT), a a a i JH III RNA (NITOR) a a a a a  
 a b RNA i a i a TOR a a i JH bi i a AA - a V i i  
*N. lugens* (L., 2016). O a NITOR a i a i a a a .NITOR a -  
 a i i a a i i NIE74Ba NIEcR.T a i i a TOR a a ,  
 JH bi i a a i a a a b i .  
 I , NITOR a i NIFoxA a NIVg.T i a -  
 a i Z ai a. (2015). W a i a TOR i a i i i BPH a i  
 , a i i a i i i a a b a i a b a i i  
 NIFoxA a NIVg.

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T i a b a a Na i a Na a S i F a i C i a (N .31171846).

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