

LARGE-SCALE BIOLOGY ARTICLE

G m -W Ma U ca a C a6 Ta c
R G a a R N c a mRNA Ca -B C m
C a a a RNA D ca A ab [OPEN]

X a, Y ,^{a,1} Ma R. W. ma,^{a,1,2} S J. A. ,^{a,b} a, B a, D. G a,b,3

D B 19104
B 19104

C D D : 0000-0002-5730-8802 (. .); 0000-0002-5782-7303 (. .A); 0000-0001-7532-0138 (B.D. .)

INTRODUCTION

¹ ; , C. ,
²C. , C. ,
14853.

³A @ fi

A () : B D.

OPEN A / / / 10.1105/ 16.00456

RESULTS

G M U C T I U v S u m RNA-D c Ta RNA
 C a G a a C a a a a RNA D ca
 P a T a c m

A- fl. C
 A fl. A 5'
 A , (A) 2008;
 , 2014). fi A ,
 (A 10 F 1A). A
 79 82% 5' (A)- A
 C , 100-
 C . F , C
 (= 0.93 (F 2A), C
 . A ~80% 5' A
 (CD) ~10% 3'
 (), <5% 5' (F 2B).
 3' A , 5'

5' A 16 17 5'
 A (~30)
 A A (2015; , 2013). A- A
 A (D , 2011).
 A C F. 1B). A , A
 A 5' 16 17 A , A ,
 5' (F. 1C) (, 2015).
GMUCT R 6 a m RNA-M a Ta RNA
C a6a E fic c
 A 45 A
 A B (,
 2012). C
 A fl 410 A 100 fl 2-
 C fl 8- 100 fl 2-
 fl (fi) 5'
 F. 1B D 1). fi ,
 fi fi (< 2.2×10^{-100} , x^2) 5'
 100- fl (F. 1A). A
 A A
 A- fi A fi
 fi fi A fi
 (D. 1). F. 172 A , TOE2 AP2, 156
 A , SPL15 SPL3 (, 2009; A .
 , 2003; z , 2008),
 fi A fi
 (F. 1B; F. 3A 3C). fi
 396 (GRF4). fi
 (, 2014), fi (F. 1B; F. 3D
 D. 1). A- fi
 AP2 (172), SPL3 (156), CSD1 (398
), (C , 2004;
 , 2007; D B , 2008).
 A A

A-
A- fi 410 A
A- fi (D 1).

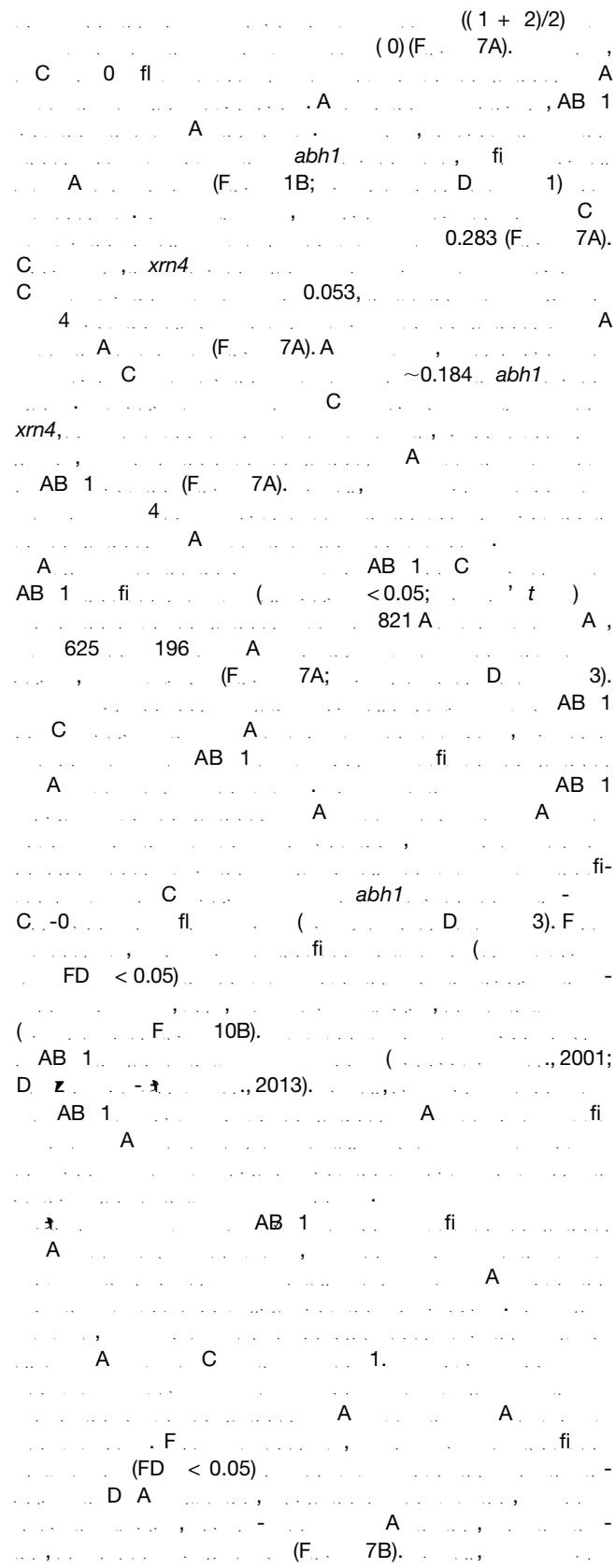
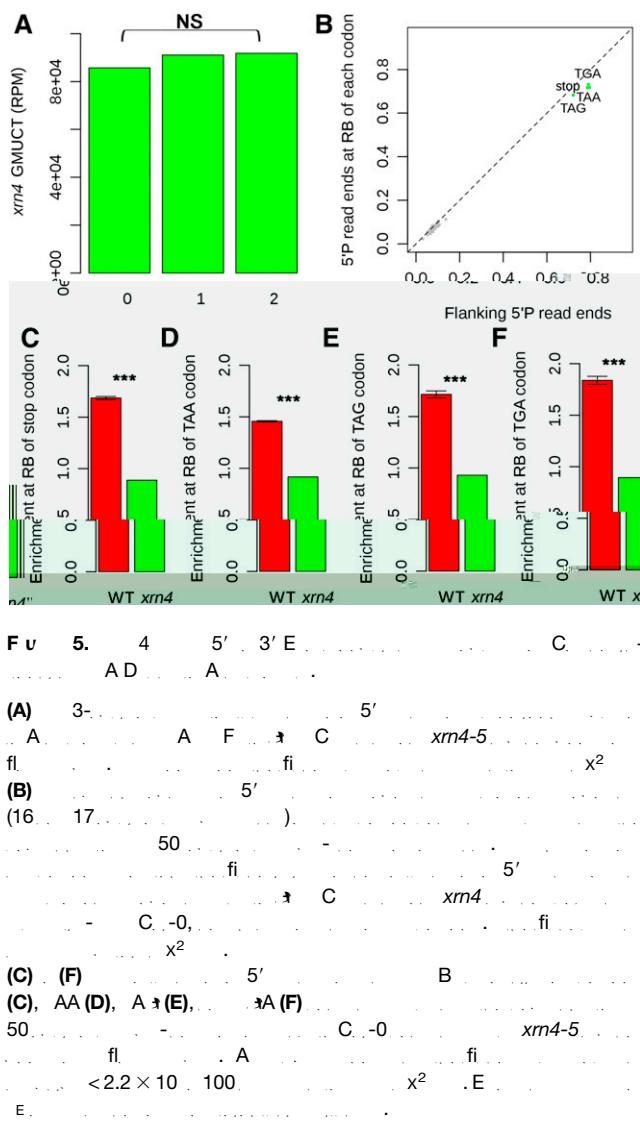
G M U C T R G a R b m a P a D T a a
T m a a a 3-N c P c mRNA O
R a F am

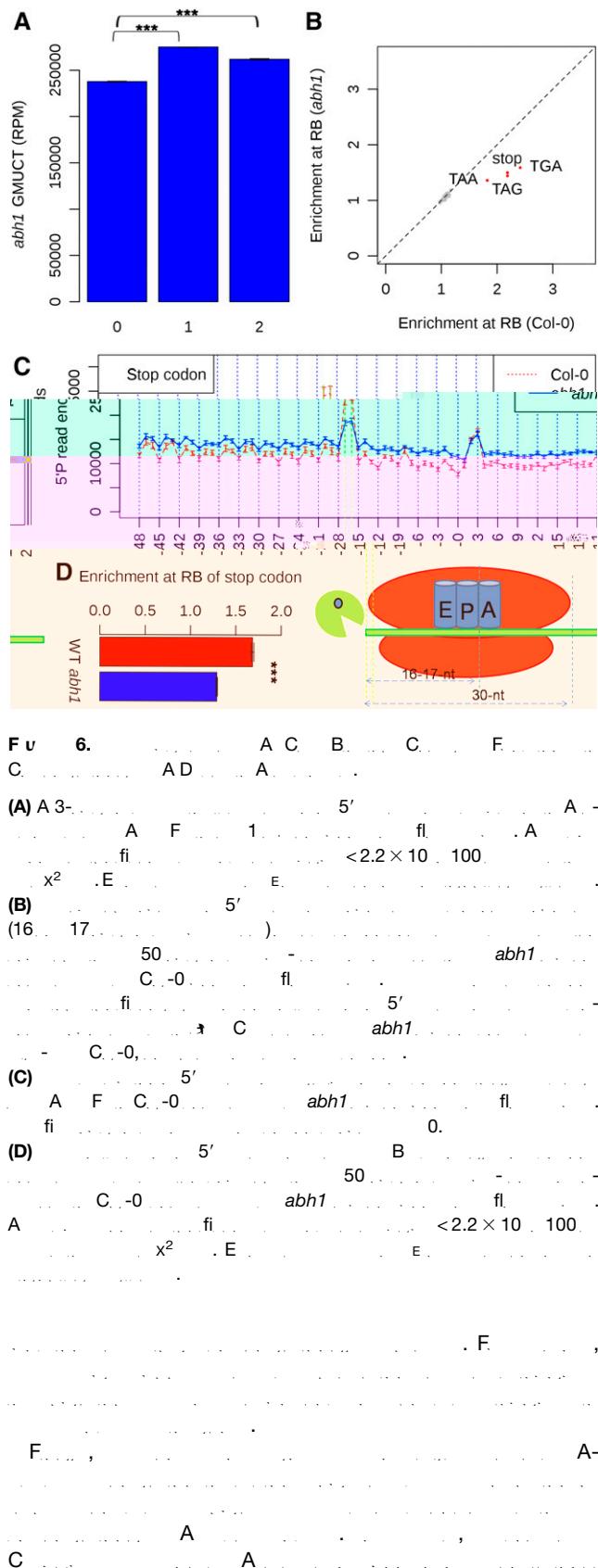
B A- A- C
A- , A- C
fi A C (F.
1B),

A C 5'
A (F 4 fi 5').

A fi ($< 2.2 \times 10^{-100}, \chi^2$) 5'
16 17 A
fi A (F 2A). A -
2000
5'
($< 2.2 \times 10^{-100}, \chi^2$). 5'
16 17 B 115 E / 38.8() 1 0 E B -0.0169 8.4682008.468253.2341184.41557 / 31 () 0 E

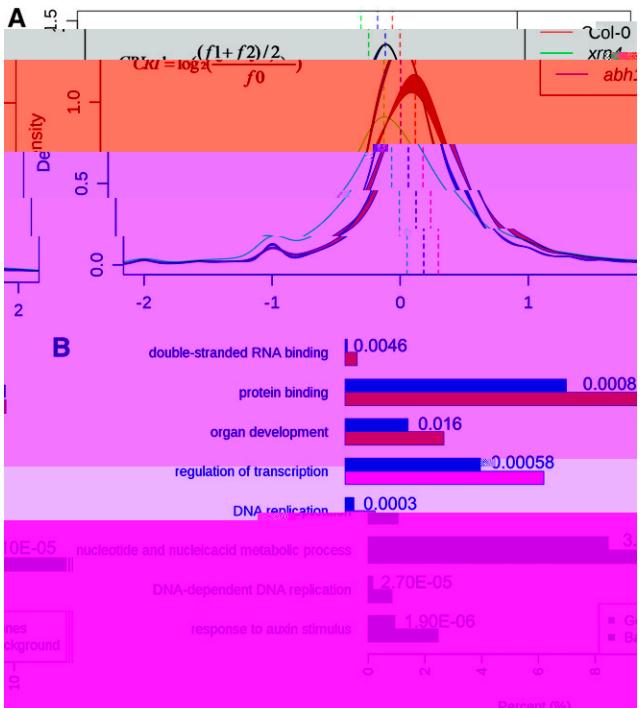
(AT4G34590) ... *XIPOTL1* (AT3G18000) (... A ... , 2014;
..., 2004; ..., 2006). A ... , ...
F ... 5'
ATCIPK6 (AT4G30960) ... *BZIP44* (AT1G75390) (...
..., 2012; ..., 2007; ..., 2013).
fi 80 F





(..., 2013) F, fl
fi (..., <0.05; χ^2) C fl
A A A fl
(..., F, 10A). A-
A .

DISCUSSION



F u 7. C AD C -0,
xrn4 abh1.

(A) C₀-0 (), xrn4 (), abh1 (); A 4

AB 1 A A

(B) A A

C () FD ().

A

A

A *S. cerevisiae* (

2015), C 5'

14 3- C 13

C -0 C (F. 4).

F , 5'

(F. 2; F. 7). , 5'

(, 2015), 1 2 0

16 17

(F. 2). fi 5'

7). (F.

16 17. AA A

.A , 5'

.A , fi

C fi (F. 6),

S. cerevisiae 3-

5'

(, 2015). fi , -

3' (F. 2B), 5' 5'

(4) A

C , fi C

CD (5'),), fi 50

(, 2011; , 2010; , 2015).

A , 3-

16- 17-

(B). 3' (F. 2A),

S. cerevisiae (, 2015).

fi

A ,

Drosophila melanogaster

(, 2011; D , 2013;

, 2014).

S. cerevisiae,

(CB 20 CB 80/AB 1), A
A , A ,
A , A ,
2008; , 2009). A

	A	.	C	
<i>abh1</i> ,				
A				fi ,
AB 1,	A			~50%
	xrn4			
	(F	5, 6,	7A).	
821	C			fi
<i>abh1</i>	C	-0		, 76%
A	fi		C	,
	A			AB 1
			C	
AB 1				
	AB 1			
	(, 2001; D
2013)				

2013), A . F . -

A , AB 1 A

AB 1 XRN4 A
 fi abh1 (2008).
 fi XRN4 abh1 A
 C -0

abh1 AB 1 (1) A ;(2) A

; (3) A A

AB 1 A
C A
A fl. A

METHODS

Pa Ma a a GMUCT L b a C u c
 C A
 5 9 4- Arabidopsis thaliana C -0 (A)
 abh1 abh1-1 abh1-8) fl.
 C 2.0 (2014). A 5' A
 (A)⁺ (A)⁺
 . A A
 3' 5'
 F C fi
 C fi

Ma GMUCT R a mRNA T a c
 A C (50-) A 10.
 A (2.4.0 " F 10-
 F 10- F 0.10" (D.
 , 2013). A fi BED fi
 fi (5') 5' E C

, 5' 100- A
 C fi
 . A , 5' 5' ,
 CD , 3' A

P c Ca c m RNA Ta a P c C a G a S
 45 A A
B B
 A (:// / A)
 21- A
 10 11 A
 5' 5' 100- fl
 A fl
 A 2 5'
 A 5'
 100- fl
 A 2 5'
 A 5'
 A 5'
 A A
 χ^2 fl
 5' A fl
 5' -

5' (0). C : $CRI = \log_2 \left(\frac{(f_1+f_2)/2}{f_0} \right)$. C
 abh1 xm4 A C C -0 C
 fi abh1 abh1 C -0 C -0
 fi t

D fi a 3-N c P c Pa ORF a E c m
 A a 5' P R a E Acc m a D R b m Pa a
 Eac C

0 (0),
 1 (1), 2 (2), 0 0
 1 2 5'
 F fi
 5' fi
 , χ^2 fl
 5' (16 17 5'
 fl 100 C -0
 C , xm4, abh1.
 abh1 fi
 , χ^2

P c B a G T a a a Ac G u ORF I fi
b GMUCT

A 5'
 F fi A , AC , C
 fi F F
 fi 5'
 2- 16 / 17
 fl 100

M au m C a a a RNA D ca

C A
 C 2
 5' (1 2)

1. C. -0... abh1
 F. B.
S m a Da a S 1. C. A.
S m a Da a S 2. F. E. 5'
 E. C. B. C.
 F.
S m a Da a S 3. A. fi... D.
 C. AD. C. -0... abh1

ACKNOWLEDGMENTS

This work was supported by grants from the National Institutes of Health (F. (CB-1243947) -
 1444490), B.D. -

AUTHOR CONTRIBUTIONS

B.D. A.
 C. B.D.
 B.D. A.

8, 2016; A., 2, 2016; 7, 2016;
 7, 2016.

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 A C . B . 18: 758–762.
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 fl A A E A A2-.
 C . 15: 2730–2741.
C a Ga, J.A., a . (2007). A
 A C . 131: 1340–1353.
C X. (2004). A A A E A A2-.
 A fl 303: 2022–2025.
C ba, Y., a , G , P. (2009). A
 B . 52: 114–124.
Da, X., a , Z a , P.X. (2011). A A
 A A 39: 155– 159.
Da a-G c, A., W a , W., R c , M., S a , I.,
 Ma , M., S a-K a, Z., a , Ja m , A.
 (2013). A 1
 :AB 1(CB 80) . AB 4- ABA .
 B . 81: 189–209.
D b A., DaG , C.A., Sc , F., D , J., Za , C.,
 J a, S., Bau , P., C a , M., a , G , a , T.R. (2013). A :
 A B 29: 15–21.
D a, D.V., a , Ba , B. (2008). A
 398, C.

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M.C., C., C., Y.Y., D., J.M., a., B.,
A., C. (2013). *Plant Cell*, **1**: 1–12.
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D., J., P., C., C., Y.Y., G., P.J., D., J.M.,
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Genome-Wide Mapping of Uncapped and Cleaved Transcripts Reveals a Role for the Nuclear mRNA Cap-Binding Complex in Cotranslational RNA Decay in Arabidopsis

M J. A B D. G
Plant Cell 2016;28:2385-2397; O 7, 2016;
DOI 10.1105/tpc.16.00456

N 11, 2016

Supplemental Data	http://www.plantcell.org/content/suppl/2016/10/17/tpc.16.00456.DC2.html http://www.plantcell.org/content/suppl/2016/10/07/tpc.16.00456.DC1.html
References	47 , 20 : :// . . / , 20 /28/10/2385. . # - -1
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