

10.1071/CP19219_AC

© CSIRO 2020

Supplementary Material: *Crop & Pasture Science*, 2020, 71, 47–55.

Functional analysis of the soybean gene *GmTIR* under biotic and abiotic stresses

Xiaoting Wang^A, Lu Huang^A, Xiaochun Bian^A, Zhan Li^A, Ruifang Gao^A, Xing Zhang^A, Xiaoli Zhang^A, Xiangnan Li^A, Haitang Wang^A, Na Guo^A, Jianying Feng^A, Jinming Zhao^{A,B} and Han Xing^{A,B}

^ASoybean Research Institute, National Center for Soybean Improvement, Key Laboratory of Biology and Genetic Improvement of Soybean (General, Ministry of Agriculture), State Key Laboratory of Crop Genetics and Germplasm Enhancement, Jiangsu Collaborative Innovation Center for Modern Crop Production, Nanjing Agricultural University, Nanjing, Jiangsu 210095, China.

^BCorresponding authors. Email: jmz3000@126.com; hanx@njau.edu.cn

Supplementary Figures

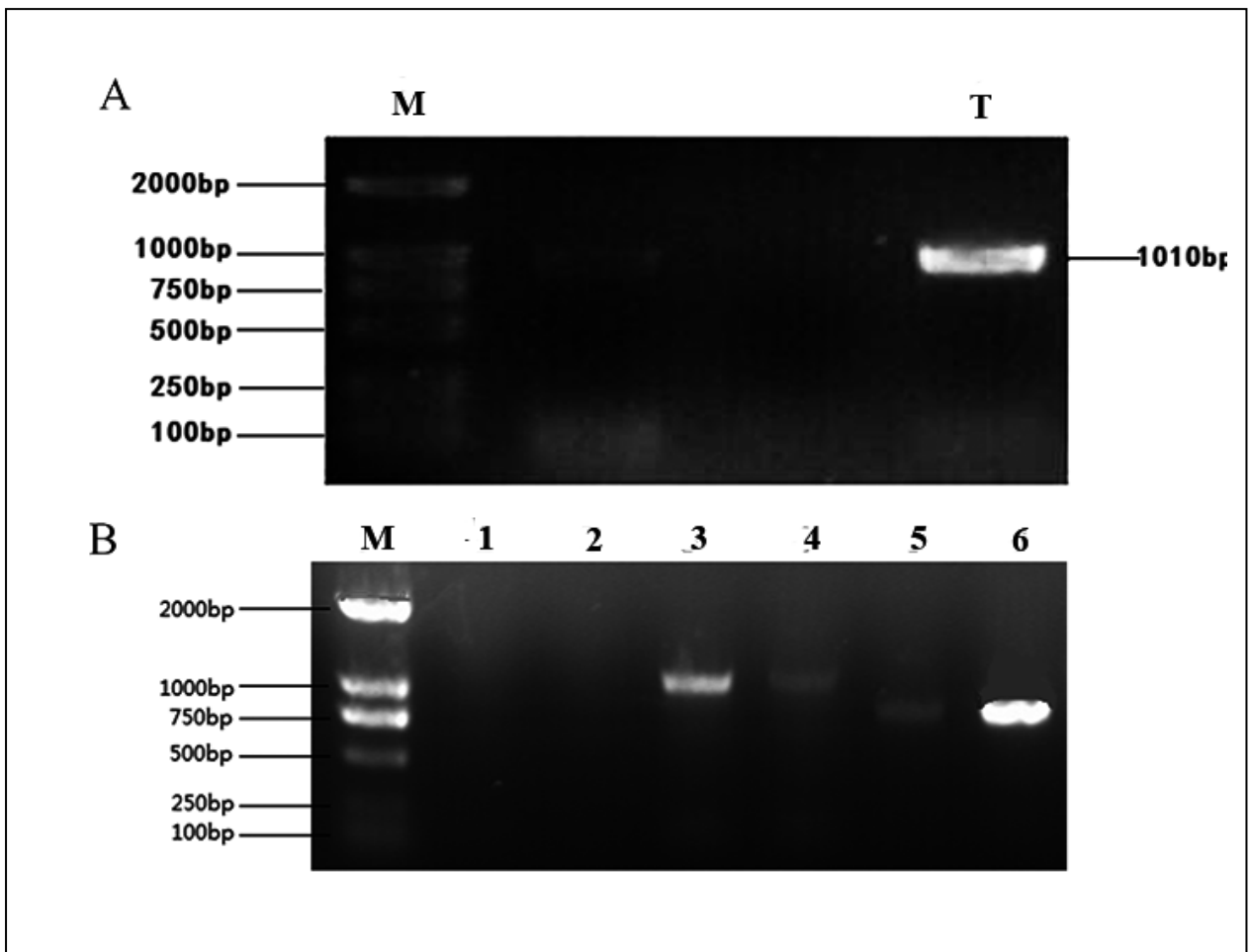


Fig. S1 PCR product electrophoresis in an agarose gel

A. Isolation of *GmTIR*

B. Construction of *GmTIR* plant expression vectors

M: DL 2000 Marker; T: PCR product of *GmTIR*

M: DL 2000 Marker; 1-2: H₂O; 3-4: 35S+*GmTIR*; 5-6: *GmTIR*

1 TCTAAAATTTTCACACCTCATTAAAGCCCTTTAGCCAAGAGTCCCTAGCCAAAAAATTAAC GA
3
63 **ATGGCATGGTCAACTTCCTCCTCTAGCACCCCTCATCAAAAGCATGAAGTGTTCCCTCAGC**
M A W S T S S S S T P H Q K H E V F L S
123 **TTCAGAGGTGAGGACACACGCTACACCTTCACTGGGCACCTTCATGCCTCTTTAACAAGA**
F R G E D T R Y T F T G H L H A S L T R
183 **CACCAAGTCAACACATACATTGACTACAACCTTCAGAGAGGTGAAGAGATATCATCCTCA**
H Q V N T Y I D Y N L Q R G E E I S S S
243 **CTTCTTAGGGCCATTGAGGAGCAAACTCTCAGTCGTTGTTTTCTCCAAAACTATGGA**
L L R A I E E A K L S V V V F S K N Y G
303 **AATCCAAGTGGTGTCTTGATGAGCTTCTGAAAATACTTGAATGTAAGAATATGAGGGGG**
N S K W C L D E L L K I L E C K N M R G
363 **CAGATTGTGTGGCCAATTTTCTATGACATTGATCCCTCGCACGTGCGCAACCAGACGGGA**
Q I V W P I F Y D I D P S H V R N Q T G
423 **ACGTATGCTGAGGCATTTGCTAAAACATGAGAAACACTTGCAGGGTCAAATGGTCAAGGTG**
T Y A E A F A K H E K H L Q G Q M V K V
483 **CAAAAGTGGAGGGTGGCTCTCAGAGAAGCAGCTAATCTCTCTGGTTGGGATTGCTCTGTC**
Q K W R V A L R E A A N L S G W D C S V
543 **AACAGGATGGAGTCTGAACTAATCGAGAAAATTGCAAAGGATGTATTAGAAAACTAAAT**
N R M E S E L I E K I A K D V L E K L N
603 **CGAGTATACGTTGGCGACCTAGATCAGCAGATTGCAAAATTAGAGCAACTTGCACAGCTT**
R V Y V G D L D Q Q I A K L E Q L A Q L
663 **CAGCATCAATTTTGCAAAACATACCTTCTTTGGAAAATGTGAGAAATCATAGAGCAACT**
Q H Q F L Q N I P S L E N V R N H R A T
723 **G TTCAGCGTATCACGGA ACTTAAAATGGAGAGAAGTGTC CGCATGCTTCGTCTCCCTCCC**
V Q R I T E L K M E R S V R M L R L P P
783 **GACATGCTTTCACATTTGGAGAATTCAAACAATAATAATTATTTTCCTTTG**TAAT**GTGCA**
D M L S H L E N S N N N N Y F P L *
843 **AAGAAATTAATTGTTGTTGTCACAAGTTTTAATGAGATACGATTCTTAATTTCAAAGTC**
903 **CCTTGTAGTTTATTTTGTATTGAAACCTTTTGGAAAATTTATGTCC**

Fig. S2 The ORF and amino acid sequences of *GmTIR*

Initiation and termination codon are shown in boldface □ the TIR domain is underlined