### **METHODOLOGY**



## A protoplast-based transient gene expression assay for the identification of heat and oxidative stress-regulatory genes in perennial ryegrass

Shanshan Lei<sup>1</sup>, Yaolong Zhu<sup>1</sup>, Weiyu Jia<sup>1</sup>, Jing Zhang<sup>1</sup>, Yingjun Chi<sup>1</sup> and Bin Xu<sup>1\*</sup>

### Abstract

**Background** With the accumulating omics data, an efficient and time-saving transient assay to express target genes is desired. Mesophyll protoplasts, maintaining most stress-physiological responses and cellular activities as intact plants, offer an alternative transient assay to study target genes' effects on heat and oxidative stress responses.

**Results** In this study, a perennial ryegrass (*Lolium perenne* L.) mesophyll protoplast-based assay was established to effectively over- or down-regulate target genes. The relative expression levels of the target genes could be quantified using RT-qPCR, and the effects of heat and  $H_2O_2$ -induced oxidative stress on protoplasts' viability could be quantitatively measured. The practicality of the assay was demonstrated by identifying the potential thermos-sensor genes *LpTT3.1/LpTT3.2* in ryegrass that over-expressing these genes significantly altered protoplasts' viability rates after heat stress.

**Conclusion** This protoplast-based rapid stress regulatory gene identification assay was briefed as 'PRIDA' that will complement the stable genetic transformation studies to rapidly identify candidate stress-regulatory genes in perennial ryegrass and other grass species.

Keywords Ryegrass, Protoplast, Oxidative stress, Heat stress, Transient gene expression

### Introduction

With increasingly accumulated omics data, time-saving assays for rapid identification of target genes are highly desirable. To identify and characterize candidate stress-regulatory genes, over-expressing or knocking out/ down the target genes is required. Yet, generating stable transgenic plants can be both time-consuming and challenging for most non-model species. Transient assays, such as virus-induced gene silencing (VIGS) [12],

*Agrobacterium*-inoculation or Agroinfiltration [3, 6], and protoplast transfection [18], were developed to complement the method of stable genetic transformation to identify or narrow down the list of candidate genes in a short while.

Mesophyll protoplasts are not only amenable to DNA transfection but also maintain most stress-physiological responses and cellular activities as intact plants [13]. Therefore, uniform and abundant isolated protoplasts offer a versatile cell-based experimental system to investigate protein subcellular localization, protein–protein interaction, cell-autonomous regulation, and oxidative stress- and innate immune responses [1, 5, 15, 18].

Perennial ryegrass (*Lolium perenne* L.) is an important perennial cool-season grass widely used for turf



© The Author(s) 2024. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/. The Creative Commons Public Domain Dedication waiver (http://creativecommons.org/publicdomain/zero/1.0/) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

<sup>\*</sup>Correspondence:

Bin Xu

binxu@njau.edu.cn

<sup>&</sup>lt;sup>1</sup> College of Agro-Grassland Science, Nanjing Agricultural University, Nanjing 210095, People's Republic of China

Solutions	Components	Note
Enzymatic hydrolysate	0.6 M D-Mannitol	Dissolve in $dH_2O$ with a final volume of 20 mL and filter through the 75 $\mu m$ nylon mesh. Freshly prepared before use
	10 mM MES·H <sub>2</sub> O	
	0.3 g Cellulase Onozuka R-10	
	0.15 g Macerozyme R-10	
W5 solution	154 mM NaCl	Dissolve in dH $_2\text{O}$ with a final volume of 200 mL and filter through the 75 $\mu\text{m}$ nylon mesh. Freshly prepared before use
	2 mM MES·H <sub>2</sub> O	
	125 mM CaCl <sub>2</sub>	
	5 mM KCl	
MMG solution	0.6 M D-Mannitol	Dissolve in $dH_2\text{O}$ with a final volume of 20 mL and filter through the 75 $\mu\text{m}$ nylon mesh. Freshly prepared before use
	4 mM MES·H <sub>2</sub> O	
	15 mM MgCl <sub>2</sub>	

Table 1 Solutions for ryegrass mesophyll protoplast isolation and transfection

and forage purposes. Stable genetic transfection of the grass takes at least 7 months from callus initiation to the regeneration of transgenic plants [4]. Ryegrass leaves covered with shining thick cuticles are not amenable to Agroinfiltration or biolistic-based transient gene expression assays. Therefore, a mesophyll protoplast isolation and DNA transfection protocol were developed for protoplast-based transient gene expression assays [19]. This transient assay has been applied in studies on the effects of exogenous supplementation of sodium copper chlorophyllin on  $H_2O_2$ -induced oxidative stress tolerance [20] and on a target gene's (*LpCYP72A15*) effect on osmotic and oxidative stress tolerance [15].

In this study, we aimed to use the ryegrass mesophyll protoplast-based transient assay to identify candidate heat and oxidative stress-regulatory genes. Using this assay, we have identified several stress-regulatory genes including the potential thermosensors in ryegrass, *Thermo-tolerance 3.1* and *3.2 (LpTT3.1 & LpTT3.2)* [22].

### **Materials and methods**

### Plant material and growth conditions

Two ryegrass varieties were used in this study, namely cv. 'Buena vista' (heat-sensitive variety) and cv. 'Xialu 6#' (heat-tolerant variety bred by our group). A healthy plant material is a prerequisite for successful protoplast isolation, transfection, and the protoplast-based gene expression assay. Perennial ryegrass plants were grown in vermiculite: perlite: peat moss (1:3:9) in a growth chamber at 25/20 °C (day/night), 14/10 light/dark photoperiod, watered twice a week, and fertilized once a week with half-strength Hoagland solution. The middle section of the first fully expanded leaf was used as the material for the isolation of protoplasts.

### Gene cloning and vector construction

Perennial ryegrass NON-YELLOW COLOURING 1-like gene (LpNOL) [21], STAYGREEN (also named as NON-YELLOWING 1, LpSGR) [17], Thermo-tolerance 3.1 of L. perenne (LpTT3.1, NCBI Genebank NO.: OR282705) and LpTT3.2 (Genebank NO.: OR282706) were used in this study. LpSGR and LpNOL were two characterized genes already known for their roles in plant heat tolerance using stable transgenic ryegrass [8, 20, 23], and the two uncharacterized genes (LpTT3.1 and LpTT3.2) were orthologous to the rice OsTT3.1 and OsTT3.2 which were known as potential thermosensors and regulators in rice [22].

To generate the RNAi hairpin structure, the selected fragments of *LpSGR* and *LpNOL* were cloned into the Sall/EcoRI and HindIII/XbaI sites pENTR D-Kannibal vector [16], and then subcloned to the pVT1629 vector [16] driven under the maize ubiquitin promoter using the LR clonase (Invitrogen). To over-express *LpTT3.1* and *LpTT3.2*, we cloned these two genes using the pENTR/D vector first (Invitrogen) and then subcloned them to the pVT1629 vector using the LR clonase as well.

### Table 2 Primers used in the study

Primer name	Sequence (5 <sup>′</sup> –3 <sup>′</sup> )
<i>LpelF4A</i> -qRT F	AACTCAACTTGAAGTGTTGGAGTG
<i>LpelF4A</i> -qRT R	AGATCTGGTCCTGGAAAGAATATG
<i>LpTT3.1-</i> qRT F	ATGCAGCGGCGGGG
<i>LpTT3.1-</i> qRT R	AACATCATATACGGGCATGCGTTC
<i>LpTT3.2-</i> qRT F <i>LpTT3.2-</i> qRT R <i>LpSGR_</i> qRT F <i>LpSGR_</i> qRT R <i>LpNOL_</i> qRT F <i>LpNOL_</i> qRT R	ATGGCTTCCTCCATCGC CAATGTTTTATCCTTCCTCTCTA GAGGAGGCGAACTCGAAG GGTTGTACCACCCTTGCAG GCTGGCAAAGAAGTTTCTCA ATGCTGCTCTCCAAATTCCT

### Protoplast isolation and transfection

The procedure of ryegrass mesophyll protoplast isolation and transfection was described [19]. In brief, the first fully expanded leaves were taken, washed twice, and then blotted dry with absorbent paper. The middle parts of the leaves (~0.5 g) were minced into fine fragments about 0.5-1 mm wide in parallel to the leaf vein in a Petri dish containing the enzymatic hydrolysate. Then, the Petri dishes were vacuumed for 30 min using a vacuum pump (60L min<sup>-1</sup> pumping rate, Product model: GM-1.0A, Jinteng Co., TianJin, China), and then gently rotated on a horizontal shaker for enzyme digestion for 4 h (h) at room temperature. The enzymatic hydrolysate was filtered through a sterilized 75 µm nylon mesh in a 50 ml centrifuge tube with  $100 \times g$  centrifugation force for 3 min at room temperature. The supernatant was gently removed with a pipette. To each 50 mL centrifuge tube, 20 mL of sterilized W5 solution was added to gently resuspend the protoplasts. Then, the protoplasts were precipitated with centrifugation at  $100 \times g$  for 3 min, resuspended with 15 mL sterilized MMG solution, and then centrifuged for another 3 min. After removing the supernatant, the precipitated protoplasts were resuspended in 2 ml of MMG solution and kept on ice before transfection. The density of protoplasts ( $\sim 5 \times 10^5$  cells per ml) was counted with a hemocytometer under a light microscope (Olympus Model BX53, Tokyo, Japan).

Then 10 µg plasmids and 100 µl protoplasts were added into 1.5 ml centrifuge tubes (the tubes were already rinsed with 1% BSA solution) and gently mixed; then 110 µl of pre-warmed PEG4000 solution (42 °C) was added to each centrifuge tube, mixed gently to a homogeneous liquid phase, and stood at room temperature for 20 min. Then, 800 µl of W5 solution was slowly added along the tube wall to each centrifuge tube, gently mixed to a homogeneous liquid phase, centrifuged at  $100 \times g$ for 3 min, and resuspended with 800 µl W5 solution. The protoplast integrity was detected again under a light microscope. The transfected protoplasts were placed in a 25 °C incubator for 16 h to allow the transgenes' expression [19].

### **Reagents and solutions**

The formula and preparation procedures of reagents and solutions are listed in Table 1 according to Yu et al. [19].



**Fig. 1** Transient suppression of *LpNOL* or *LpSGR* altered heat and oxidative stress tolerances of ryegrass protoplasts (cv. 'Buena Vista'). **A** Ryegrass protoplasts transfected with pVT1629-GFP visualized under bright (left) and GFP (right) fields. Bars in (**A**) indicate 1 mm. **B**, **C** Relative expressions of *LpSGR* and *LpNOL* after 16 h of protoplasts transfection with the empty vector control and RNAi vectors. **D** Protoplasts' viability after heat stress at 35 °C for 20 min. **E** Protoplasts' viability under 25–50 mM H<sub>2</sub>O<sub>2</sub>-induced oxidative stress. Noting that the control in (**B**–**E**) was those transfected with empty vectors. At least 30 protoplasts were counted in one optical field, and the means were from 3 optical fields. Different letters represent statistically significant differences at *P*=0.05, and the bars above the columns represent standard errors

Heat stress and  $H_2O_2$  treatments of transfected protoplasts After 16 h of incubation, the transfected protoplasts were ready for heat and oxidative stress treatments. For heat stress, tubes containing protoplasts were transferred to 35 °C or at 31 °C, 33 °C, 35 °C, 37 °C, 39 °C, or 41 °C for temperature range test, in water baths for 20 min, stained with 1 µl of 0.1% Evans blue dye for 1 min and then observed under microscope (×100). Three representative optical fields were photographed.

For  $H_2O_2$ -induced oxidative stress, 30%  $H_2O_2$  solution was added to the suspended protoplasts in a centrifuge tube to make the final  $H_2O_2$  concentration of 25, 37.5, or 50 mM, incubating for 5 min at room temperature (note: To make 1000 mM  $H_2O_2$  stock solution, add 10.21 ml 30%  $H_2O_2$  solution to 89.79 ml water. The  $H_2O_2$  solution shall be freshly made before use). Then, 1 µL of 0.1% Evan blue solution was added and gently mixed, incubating for 1 min, and then immediately photographed at the microscope as described above. Broken protoplasts stained in blue were counted as dead protoplasts and the number of viable and total protoplasts was counted. Protoplast viability (%) was counted as the number of intact, unstained protoplasts divided by the number of total protoplasts.

For the detection of relative expression levels of *LpTT3.1* and *LpTT3.2* in ryegrass after heat stress, potted ryegrass plants (cv. 'Buena vista' and cv. 'Xialu 6#') were treated under 38 °C for 24 h (h) in a growth chamber. The fully expanded leaves were harvested from the plants for RNA extraction.

### RNA extraction and RT-qPCR

The MolPure<sup>®</sup> cell RNA kit (Yeasen Biotech. Co., Ltd., Shanghai, China) was used for RNA extraction from protoplasts, and the MolPure<sup>®</sup> Plant RNA Kit (Yeasen Biotech. Co) was used for RNA extraction from the whole leaves. The quantity and integrity of total RNA were



**Fig. 2** Sequence alignments and heat-responsive expression of *LpTT3.1 & LpTT3.2*. **A**, **B** Sequence alignment of LpTT3.1 vs OsTT3.1 and LpTT3.2 vs OsTT3.2; **C**, **D** Relative expression of *LpTT3.1 & LpTT3.2* after heat stress at 38 °C in two varieties of perennial ryegrass using RT-qPCR. Different letters represent statistically significant differences at P = 0.05, and the bars above the columns represent standard errors



**Fig. 3** Transient over-expression of *LpTT3.1* or *LpTT3.2* altered heat and oxidative stress tolerances of ryegrass protoplasts (cv. 'Buena Vista'). **A**, **B** Relative expression levels of *LpTT3.1* and *LpTT3.2* after 16 h of protoplast transfection with the empty vector control or target genes under a *ubiquitin* promoter; **C** Protoplast viability after heat stress at 35 °C for 20 min; **E**, **F** Protoplasts' viabilities under heat stress at 31–41°C for 20 min or under 25–50 mM H<sub>2</sub>O<sub>2</sub>-induced oxidative stress. Noting that the control in (**C**–**F**) was those transfected with empty vectors. 'ND' in (**G**) is the abbreviation for 'not detected'. At least 30 protoplasts were counted in one optical field, and the means were from 3 optical fields. Different letters represent statistically significant differences at P=0.05, and the bars above the columns represent standard errors

checked using Nanoready (NanoDrop Technologies, Hangzhou, China) and by running through a 1% agarose gel. The cDNA synthesis was performed using the ChamQ Universal SYBR qPCR Master Mix for RT-PCR Kit (Vazyme Biotech Co., Nanjing, China) with oligo-dT primers. For reverse transcription-quantitative polymerase chain reaction (RT-qPCR), PRIMER EXPRESS\_ software (v3.0; Applied Biosystems, Foster City, CA, USA) was used to design primer sets for *LpTT3.1*, *LpTT3.2*, *LpSGR*, *LpNOL* and the reference gene (*LpeIF4A*) [7] (Table 2). Each sample had three replicates, and the data were normalized against the reference gene *LpeIF4A*.

### Statistical analysis

Data were statistically analyzed by One-way ANOVA at a significance level of 0.05 using SPSS20.0 software (SPSS Inc.). Data are expressed as means $\pm$ standard error. At least two separate experiments were carried out for each test.

### Results

# Suppression of *LpNOL* or *LpSGR* improved or compromised heat and oxidative stress tolerance in mesophyll protoplasts

More than 90% of the mesophyll protoplasts were successfully transfected with plasmids expressing the GFP



**Fig. 4** Transient gene suppression/over-expression of *LpNOL*, *LpSGR*, *LpTT3.1*, or *LpTT3.2* altered heat and oxidative stress tolerances of ryegrass protoplasts (cv. 'Xialu 6#'). Protoplasts' viabilities under heat stress at  $37^{\circ}$ C for 20 min (**A**) or under H<sub>2</sub>O<sub>2</sub>-induced oxidative stress (**B**). Noting that the control was those transfected with empty vectors. At least 30 protoplasts were counted in one optical field, and the means were from 3 optical fields. Different letters represent statistically significant differences at *P*=0.05, and the bars above the columns represent standard errors

reporter gene (Fig. 1A). To test whether the transfected protoplasts was a reliable and sensitive system for characterizing heat-stress regulatory genes, we transfected ryegrass (cv. 'Buena Vista') protoplasts with the RNAi vectors to suppress NON-YELLOW COLOURING 1-like (LpNOL) and STAYGREEN (LpSGR), respectively. Stable transgenic ryegrass with suppressed LpNOL or LpSGR have significantly improved or compromised heat tolerance [8, 20, 23]. After DNA transfection, the protoplasts were rested in 25 °C for 16 h, and the relative expression levels of LpNOL and LpSGR were ~ 20% that of the control ones (Fig. 1B, C). After the resting stage, the protoplasts were subjected to heat treatment (35 °C) for 20 min, then stained with Evans blue for 1 min and immediately observed under the microscope. Consistent with the results of stable transgenic ryegrass plants, transient suppression of LpNOL led to 16% higher protoplasts viability, while suppression of *LpSGR* led to 49% lower protoplasts viability than the empty vector control (Fig. 1D).

Oxidative stress is often associated with heat stress on plants with the burst and accumulation of reactive oxygen species (ROS) aggravating cellular damage [9]. Constitutive suppression of *LpNOL* and *LpSGR* also affected the ROS accumulation and the ROS-scavenging system in stable transgenic plants [8, 20, 23]. To test whether this protoplast-based transient assay is also sensitive to oxidative stress treatment, we treated the transfected protoplasts with different concentrations of  $H_2O_2$ . As shown in Fig. 1E, with  $H_2O_2$  at final concentrations of 37.5 to 50 mM, transient suppression of *LpSGR* significantly decreased protoplast viability, while the contrary was true for *LpNOL* (Fig. 1E).

### Identification of potential thermosensors, *LpTT3.1* and *LpTT3.2*, in ryegrass

To demonstrate this mesophyll-based transient expression system is suitable for identifying candidate stressregulatory genes, LpTT3.1 and LpTT3.2 were chosen as candidate genes for characterization. These two ryegrass genes were orthologous to the rice TT3.1 and TT3.2 sharing 86.4% and 74.4% nucleotide (nt) identities, respectively (Fig. 2A, B). Expression levels of both LpTT3.1 and LpTT3.2 increased after 2 h of heat stress, the expression of LpTT3.1 was more sharply increased in the heat-tolerant variety (cv. 'Xialu 6#') than in the sensitive one (cv. 'Buena Vista') (Fig. 2C, D).

Using the protoplasts of cv. 'Buena Vista', transient over-expressing LpTT3.1 resulted in 58 times higher expression levels and led to 29% higher protoplast viability rate after heat stress treatment than those of the control (Fig. 3A, C); while transient over-expressing LpTT3.2 resulted in 82 times higher expression levels but 24% less protoplast viability rate after heat stress (Fig. 3B, C). To optimize the temperature range for heat stress, we treated protoplast over-expressing  $LpTT3.1 \approx LpTT3.2$  at 31 to 41 °C, and the result showed that 35 to 39 °C was the optimal treatment condition (Fig. 3D, E). Furthermore, over-expressing LpTT3.2 significantly decreased protoplast viability when treated with 25 to 50 mM H<sub>2</sub>O<sub>2</sub>, while over-expressing LpTT3.1 did not affect protoplast viability under oxidative stress (Fig. 3F).

To verify whether the above-selected stress conditions were effective for other ryegrass varieties, we tested it in a more heat-tolerant variety (cv. 'Xialu 6#'). As shown in Fig. 4, the viabilities of transfected protoplasts after heat treatment at 37 °C or after 37.5 to 50 mM  $H_2O_2$ 



1. Protoplast preparation

Note: Protoplasts density is ~5×10<sup>5</sup> cells per ml, and their viabilities are no less than 80%.

### 7 2. Protoplast transfection



Protoplast transfection rate is no less than 50%; over 90% transfection rate can be achieved. After DNA transfection, rest protoplasts for 16 hours to allow target genes' over-expression or suppression.



3. Target gene's expression level analysis Note:

The RNA extraction shall be compatible with the suspension cell culture. Noting that regular plant RNA extraction kit may not work well for protoplasts.



#### 5. Protoplast viability analysis 70 60 Note: 50 Addition of Evans blue dye for longer than 1 min has little 40 or no effect on the protoplasts' 30 viability: 20 At least three optical fields 10 shall be observed for statistical analysis. ٥ RNAi::LpSGR RNAi::LpNOL Control



treatment differed according to the target genes, and the results were consistent with those obtained with ryegrass cv. 'Buena Vista'.

### Discussion

To explore the post-genomics data, time-saving and high throughput functional genomic assays are indispensable to identify target genes. After the release of perennial ryegrass genome [2, 10], multiple comparative transcriptome analyses and Genome-Wide Association Studies (GWAS) were carried out [11, 14, 24], providing numerous candidate genes for further functional studies. Generation of stable transgenic ryegrass plants takes at least 7 months, and more time is needed to characterize these transgenic plants for heat and/ or oxidative tolerance. To complement the extremely time-consuming stable genetic transformation procedure, here we report an alternative assay using the mesophyll protoplast-based transient gene expression system for rapid identification of stress regulatory genes in ryegrass. Therefore, we named this protocol as 'PRIDA' for Protoplast-based Rapid stress regulatory gene Identification Assay. Compared with ectopic expression in yeast or in Arabidopsis, the PRIDA assay is carried out in the native plant species and takes even less time.

Leaves of perennial ryegrass are narrow (3-5 mm width) with parallel veins and shining cuticles that are recalcitrant to Agrobacterium- or biolistic-based transient gene expression assays. Yet, abundant mesophyll protoplasts can be isolated from its green leaves, the protoplast DNA transfection rate can be higher than 90%, and these freshly isolated protoplasts maintain most of their innate cellular activities and are sensitive to abiotic stress conditions [1, 5, 15, 18]. All these features make it a feasible system to characterize the target gene's regulatory roles in stress. In a recently reported study, we used this PRIDA assay to analyze the target gene's (*LpCYP72A15*) effect on osmotic/dehydration stress [15]. In this study, we further showed that this PRIDA assay, capable of both over-expressing and suppressing the target gene, was an effective way to identify genes involved in heat and oxidative stress. A similar assay can be developed in other non-model grass species as well for the identification and characterization of genes regulating heat, oxidative, osmotic, and other types of abiotic stresses.

Fine-tuning the stress treatment conditions is a key to the successful interpretation of the results of PRIDA. Perennial ryegrass is a cool-season grass species that is sensitive to heat stress [24]. Yet, treating ryegrass mesophyll protoplasts below 33 °C cannot tell the stress difference in a short time, while treatment at 41 °C for 20 min made the protoplasts stick to each other, making it impossible to count precisely. Therefore, the optional temperature range for heat treatment is 35 to 39 °C. Treatment with 100 mM H<sub>2</sub>O<sub>2</sub> killed all protoplasts within 5 min, while treatment with 25 mM  $H_2O_2$  could not discriminate the transgenes' effects. H<sub>2</sub>O<sub>2</sub> treatment at 37.5-50 mM is the optimal condition. Similarly, mannitol at 100 mM was the optimal condition for osmotic stress treatment in this ryegrass protoplast transient assay [15]. We tested the protocol in two ryegrass varieties with contrasting heat tolerance and the results indicated that the above-selected stress conditions were optimal for both varieties. Yet, when the 'PRIDA' system is applied to other grass species, these treatment conditions shall be optimized accordingly.

The TT3.1 and TT3.2 genes act as thermosensors in rice that, upon heat stress, both the plasma membranelocalized E3 ligase TT3.1 and the chloroplast-localized TT3.2 proteins translocate to endosomes where TT3.1 trigger the degradation of TT3.2 to protect chloroplast from heat-induced damage [22]. Over-expressing TT3.1 or TT3.2 in stable transgenic rice improved or compromised plant heat tolerance, respectively [22]. In this study, using this PRIDA assay, we were able to pin down that this pair of ryegrass genes (LpTT3.1 & LpTT3.2) that likely share the same functions in heat tolerance and can be used as gene editing targets for further molecular breeding of perennial ryegrass.

It should also be noted that these mesophyll protoplasts lack integral cell wall and cannot be kept for more than a week. Therefore, this transient assay cannot be used to study target genes involving cell wall and/or inter-cellular stress-related pathways or their long-term effects on cell division, growth, and development. An in-silico analysis of the target gene/protein (e.g. prediction of target proteins' subcellular localization) shall be performed before the test.

### Conclusion

A detailed procedure of PRIDA was summarized and illustrated in Fig. 5 with critical points pinpointed. The established PRIDA assay in ryegrass can be readily adapted to other types of abiotic stresses and in other plant species to rapidly identify candidate stress-regulating genes.

### Author contributions

SL, LZ, WJ, JZ, and YC performed vector construction, ryegrass protoplast isolation and transfection experiments; SL, LZ, and BX participated in manuscript preparation; BX conceived and designed the experiments. All authors read and approved the final manuscript.

### Funding

This project was supported by National Key Research and Development Program of China (2023YFF1001403) and by a Grant from the National Natural Science Foundation of China (No. 32271755).

### Availability of data and materials

The datasets supporting the conclusions and description of a complete protocol are included within the article. The pVT1629 vector is available upon request.

### Declarations

Ethics approval and consent to participate

All authors read and approved the manuscript.

### Consent for publication

All authors agreed to publish this manuscript.

### Competing interests

The authors declare that they have no competing interests.

Received: 5 February 2024 Accepted: 24 April 2024 Published online: 09 May 2024

### References

- Asai T, Tena G, Plotnikova J, Willmann MR, Chiu WL, Gomez-Gomez L, Boller T, Ausubel FM, Sheen J. MAP kinase signalling cascade in Arabidopsis innate immunity. Nature. 2002;415(6875):977–83. https://doi.org/10. 1038/415977a.
- Byrne SL, Nagy I, Pfeifer M, Armstead I, Swain S, Studer B, Mayer K, Campbell JD, Czaban A, Hentrup S, Panitz F, Bendixen C, Hedegaard J, Caccamo M, Asp T. A synteny-based draft genome sequence of the forage grass Lolium perenne. Plant J. 2015;84(4):816–26. https://doi.org/10.1111/tpj. 13037.
- 3. Colinas M, Goossens A. Transient gene expression in *Catharanthus roseus* flower petals using agroinfiltration. Methods Mol Biol. 2022;2505:281–91. https://doi.org/10.1007/978-1-0716-2349-7\_20.
- Grogg D, Rohner M, Yates S, Manzanares C, Bull SE, Dalton S, Bosch M, Studer B, Broggini GAL. Callus induction from diverse explants and genotypes enables robust transformation of perennial ryegrass (Lolium perenne L.). Plants (Basel). 2022. https://doi.org/10.3390/plants11152054.
- Kovtun Y, Chiu W-L, Tena G, Sheen J. Functional analysis of oxidative stress-activated mitogen-activated protein kinase cascade in plants. Proc Natl Acad Sci. 2000;97(6):2940–5. https://doi.org/10.1073/pnas.97.6.2940.
- Krenek P, Samajova O, Luptovciak I, Doskocilova A, Komis G, Samaj J. Transient plant transformation mediated by *Agrobacterium tumefaciens*: principles, methods and applications. Biotechnol Adv. 2015;33(6 Pt 2):1024–42. https://doi.org/10.1016/j.biotechadv.2015.03.012.
- Huang L, Yan H, Jiang X, Yin G, Zhang X, Qi X, Zhang Y, Yan Y, Ma X, Peng Y. Identification of candidate reference genes in perennial ryegrass for quantitative RT-PCR under various abiotic stress conditions. PLoS ONE. 2014;9: e93724. https://doi.org/10.1371/journal.pone.0093724.
- Lei S, Yu G, Rossi S, Yu J, Huang B. LpNOL-knockdown suppression of heat-induced leaf senescence in perennial ryegrass involving regulation of amino acid and organic acid metabolism. Physiol Plant. 2021;173(4):1979–91. https://doi.org/10.1111/ppl.13541.
- Nadarajah KK. ROS homeostasis in abiotic stress tolerance in plants. Int J Mol Sci. 2020. https://doi.org/10.3390/ijms21155208.
- Nagy I, Veeckman E, Liu C, Bel MV, Vandepoele K, Jensen CS, Ruttink T, Asp T. Chromosome-scale assembly and annotation of the perennial ryegrass genome. BMC Genomics. 2022;23(1):505. https://doi.org/10.1186/ s12864-022-08697-0.
- 11. Nie G, Zhou J, Jiang Y, He J, Wang Y, Liao Z, Appiah C, Li D, Feng G, Huang L, Wang X, Zhang X. Transcriptome characterization of candidate

genes for heat tolerance in perennial ryegrass after exogenous methyl jasmonate application. BMC Plant Biol. 2022;22(1):68. https://doi.org/10. 1186/s12870-021-03412-9.

- 12. Rössner C, Lotz D, Becker A. VIGS goes viral: how VIGS transforms our understanding of plant science. Annu Rev Plant Biol. 2022;73:703–28. https://doi.org/10.1146/annurev-arplant-102820-020542.
- 13. Sheen J. Signal transduction in maize and Arabidopsis mesophyll protoplasts. Plant Physiol. 2001;127(4):1466–75.
- Wang K, Liu Y, Tian J, Huang K, Shi T, Dai X, Zhang W. Transcriptional profiling and identification of heat-responsive genes in perennial ryegrass by RNA-sequencing. Front Plant Sci. 2017;8:1032. https://doi.org/10.3389/ fpls.2017.01032.
- Xing J, Yang Y, Zhang Q, Yin T, Zhao R, Hao G, Liu X, Chi Y, Zhang J. A cytochrome P450 gene, LpCYP72A15, confers drought tolerance in perennial ryegrass. Grass Forage Sci. 2024. https://doi.org/10.1111/gfs.12642.
- Xu B, Escamilla-Treviño LL, Sathitsuksanoh N, Shen Z, Shen H, Zhang YH, Dixon RA, Zhao B. Silencing of 4-coumarate:coenzyme A ligase in switchgrass leads to reduced lignin content and improved fermentable sugar yields for biofuel production. New Phytol. 2011;192(3):611–25. https://doi. org/10.1111/j.1469-8137.2011.03830.x.
- Xu B, Yu G, Li H, Xie Z, Wen W, Zhang J, Huang B. Knockdown of STAY-GREEN in perennial ryegrass (*Lolium perenne* L.) leads to transcriptomic alterations related to suppressed leaf senescence and improved forage quality. Plant Cell Physiol. 2019;60(1):202–12. https://doi.org/10.1093/ pcp/pcy203.
- Yoo SD, Cho YH, Sheen J. Arabidopsis mesophyll protoplasts: a versatile cell system for transient gene expression analysis. Nat Protoc. 2007;2(7):1565–72. https://doi.org/10.1038/nprot.2007.199.
- Yu G, Cheng Q, Xie Z, Xu B, Huang B, Zhao B. An efficient protocol for perennial ryegrass mesophyll protoplast isolation and transformation, and its application on interaction study between LpNOL and LpNYC1. Plant Methods. 2017;13:46. https://doi.org/10.1186/s13007-017-0196-0.
- Yu G, Xie Z, Chen W, Xu B, Huang B. Knock down of NON-YELLOW COL-OURING 1-like gene or chlorophyllin application enhanced chlorophyll accumulation with antioxidant roles in suppressing heat-induced leaf senescence in perennial ryegrass. J Exp Bot. 2022;73(1):429–44. https:// doi.org/10.1093/jxb/erab426.
- Yu G, Xie Z, Zhang J, Lei S, Lin W, Xu B, Huang B. NOL-mediated functional stay-green traits in perennial ryegrass (*Lolium perenne* L.) involving multifaceted molecular factors and metabolic pathways regulating leaf senescence. Plant J. 2021;106(5):1219–32. https://doi.org/10.1111/tpj. 15204.
- Zhang H, Zhou JF, Kan Y, Shan JX, Ye WW, Dong NQ, Guo T, Xiang YH, Yang YB, Li YC, Zhao HY, Yu HX, Lu ZQ, Guo SQ, Lei JJ, Liao B, Mu XR, Cao YJ, Yu JJ, Lin Y, Lin HX. A genetic module at one locus in rice protects chloroplasts to enhance thermotolerance. Science. 2022;376(6599):1293–300. https://doi.org/10.1126/science.abo5721.
- Zhang J, Li H, Huang X, Xing J, Yao J, Yin T, Jiang J, Wang P, Xu B. STAYGREEN-mediated chlorophyll a catabolism is critical for photosystem stability during heat-induced leaf senescence in perennial ryegrass. Plant Cell Environ. 2022;45(5):1412–27. https://doi.org/10.1111/pce.14296.
- Zhang J, Li H, Jiang Y, Li H, Zhang Z, Xu Z, Xu B, Huang B. Natural variation of physiological traits, molecular markers, and chlorophyll catabolic genes associated with heat tolerance in perennial ryegrass accessions. BMC Plant Biol. 2020;20(1):520. https://doi.org/10.1186/ s12870-020-02695-8.

### **Publisher's Note**

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.