## **Chapter 5**

### Analyzing Synthetic Promoters Using Arabidopsis Protoplasts

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#### Abstract

This chapter describes a transient protoplast co-transfection method that can be used to quantitatively study in vivo the activity and function of promoters and promoter elements (reporters), and their induction or repression by transcription factors (effectors), stresses, hormones, or metabolites. A detailed protocol for carrying out transient co-transfection assays with Arabidopsis At7 protoplasts and calculating the promoter activity is provided.

Key words Arabidopsis thaliana, At7 cell culture, Protoplast co-transfection assays, Reporter gene

#### 1 Introduction

For high-level constitutive expression, or for precise control of transgene activity in response to a specific stimulus, promoters are the key for successful genetic engineering strategies. Therefore, detailed knowledge about the concerted action of both cis- and trans-acting elements is necessary. Defining cis-acting elements and the characterization of transcription factors which bind and/ or regulate a promoter of choice is a standard experimental approach. Since in vitro DNA-binding assays might not reflect the situation in a living cell, in vivo assays are favored. The yeast onehybrid assay [1] is the method of choice for rapid detection of protein-DNA interactions. However, this experimental approach has some limitations for the analysis of plant transcription factors and promoters. Among these are different conditions inside the yeast nucleus compared to the situation in plants, additional transcriptional start sites in promoters larger than approximately 300 bps in length, and false positives due to regulatory proteins with high affinity to unspecific DNA regions [2]. For the analysis of (synthetic) plant promoters, the use of plant cell systems avoids

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most of the disadvantages of yeast systems. Furthermore, plant systems provide necessary perception and signaling systems for the study of signal-induced processes.

Transient protoplast co-transfection systems are used for the study of unbiased activity and function of promoter elements, as well as for promoter activation or repression by a single or multiple transcription factors or several stresses or hormone treatments. In this respect, the transfection assay is focused on the analysis of factors that control the activity of a given promoter, and the role of proximal 5'-upstream *cis*-regulatory elements or sequences. These elements are important and central components for accurate and refined synthetic promoter design. Transient protoplast cotransfection assays allow fast access to results, especially when compared with stable transformation. In addition, they are unaffected by position effects caused by features of the site of transgene integration and by the copy number of inserted transgenes. The drawback is that cell type specificity or developmental control of promoter activity can usually not be studied in protoplasts from cultured cells.

The protoplast assay system employs purified plasmid DNA introduced into the cells via PEG-mediated DNA uptake. Various plasmids can be introduced at the same time (co-transfection). The protoplast assay system relies on assessing the level of gene expression for an engineered promoter construction that drives the expression of a reporter gene. The level of expression of the reporter, in the case described here,  $\beta$ -glucuronidase (*uidA*, GUS) is taken as a measure for promoter activity. Thus, transient protoplast co-transfection assays have provided a wealth of information about *cis* elements required for promoter function, transcription factors, and signaling proteins that regulate expression of genes and signals regulating inducible gene expression [3–8].

#### 2 Materials

2.1 Cells, Buffers, and Solutions	1. At7 cell culture: hypocotyl-derived <i>A. thaliana</i> Columbia cell culture [9], maintained at 26 °C in darkness on a rotary shaker, weekly subcultured.
	2. <i>dam– E. coli</i> : Methylation-deficient <i>dam</i> and <i>dcm E. coli</i> strain K12 ER2925 (NEB).
	3. 1000× 2,4-Dichlorophenoxyacetic acid (2,4-D): 1 mg/mL.
	<ol> <li>MS medium: 4.3 g Murashige and Skoog Basal Salt Mixture (MS, Sigma-Aldrich), 1 mL 1000× 2,4-D, 10 mL 1000× Gamborg's Vitamin Solution (Sigma-Aldrich), 30 g sucrose. Adjust to pH 5.7 with 1 M KOH and bring to 1 L with deionized H<sub>2</sub>O. Autoclave.</li> </ol>

- Cellulase-mazerozyme solution: 1.16% (w/v) cellulase Onozuka R-10 (Serva), 0.27% mazerozyme R-10 (Serva). Solve in 240 mM CaCl<sub>2</sub>, stir cautiously until enzymes are dissolved (1–1.5 h). Pass enzyme solution through a folded filter paper, then filter sterilize.
- 6. B-5 floating medium (B5 solution): 3.1 g Gamborg's B-5 Basal Salt Mixture (Sigma-Aldrich), 1 mL 1000×2,4-D, 136 g sucrose. Adjust to pH 5.7 with 1 M NaOH and bring to 1 L with deionized H<sub>2</sub>O. Filter sterilize.
- 7. 240 mM CaCl<sub>2</sub>. Autoclave.
- PEG solution: 125 g PEG 6000, 11.8 g Ca(NO<sub>3</sub>)<sub>2</sub>×4H<sub>2</sub>O, 41 g mannitol. Adjust to pH 9 with 1 M KOH and bring to 0.5 L with deionized H<sub>2</sub>O. Filter sterilize and store in 5 mL aliquots at -20 °C.
- 9. 275 mMCa(NO<sub>3</sub>)<sub>2</sub>: Adjust to pH 6.0 with 1 M KOH. Autoclave.
- 10. 0.1 M K<sub>2</sub>HPO<sub>4</sub> and 0.1 M KH<sub>2</sub>PO<sub>4</sub>. Autoclave.
- 11. 0.1 M Potassium phosphate: Mix the appropriate volumes of 0.1 M  $K_2$ HPO<sub>4</sub> and 0.1 M KH<sub>2</sub>PO<sub>4</sub> for a desired pH of 7.0. Store at 4 °C up to 1 month.
- 12. Protein extraction buffer: 100 mM potassium phosphate, 1 mM DTT (Sigma-Aldrich). Filter sterilize and store at 4 °C.
- 13. 2×Luciferase assay stock solution: 40 mM tricine, 2.14 mM  $Mg(CO_3)_4Mg(OH)_2 \times 5H_2O$ , 5.34 mM  $MgSO_4 \times 7H_2O$ , 0.2 mM EDTA. Store at 4 °C.
- 14. Luciferase substrate solution:  $1 \times$  luciferase assay stock solution, 33.3 mM DTT (Sigma-Aldrich), 270  $\mu$ M CoA trilithium salt (Sigma-Aldrich), 470  $\mu$ M luciferin (Roche), 570  $\mu$ M ATP (Sigma-Aldrich). CoA trilithium salt and luciferin are light-sensitive, keep in the dark. Check pH and adjust to pH 7.5 if necessary. Filter sterilize and store in 5 mL aliquots at -80 °C in light-tight tubes.
- 15. 0.5 M Na<sub>2</sub>HPO<sub>4</sub> and 0.5 M NaH<sub>2</sub>PO<sub>4</sub>. Autoclave.
- 16. 0.5 M Sodium phosphate buffer: Mix the appropriate volumes of 0.5 M Na<sub>2</sub>HPO<sub>4</sub> and 0.5 M NaH<sub>2</sub>PO<sub>4</sub> for a desired pH of 7.0. Store at 4 °C up to 1 month.
- 17. GUS buffer: 50 mM sodium phosphate buffer, 1 mM EDTA pH 8.0, 0.1% (v/v) Triton X-100, 10 mM β-mercaptoethanol.
- 4-MUG substrate solution: 20 mM 4-methylumbelliferyl-β-D-glucopyranosiduronic acid (4-MUG, Sigma-Aldrich). Solve in GUS buffer. Filter sterilize and store in 15 mL aliquots at -20 °C.
- 19. MU stock solution: 10 mM 4-methylumbelliferone (MU, Sigma-Aldrich). Solve in ethanol. Store at 4 °C.

- 20. MU dilution series: Dilute the MU stock solution with GUS buffer to 0, 2.5, 5.0, 12.5, 25, 50, 100, 150, 200, and 250 μM.
- 21. BSA dilution series: Dilute a 10 mg/mL BSA stock solution with protein extraction buffer to 0, 2, 4, 8, and 16  $\mu$ g per 10  $\mu$ L buffer.
- 22. Protein assay dye reagent: Dilute the protein assay dye reagent concentrate (Bio-Rad) 1:5 with deionized H<sub>2</sub>O.

#### **2.2 Equipment** 1. Laminar flow cabinet.

- 2. 50 mL Falcon tubes.
- 3. 13 mL centrifuge tubes.
- 4. 1.5 mL reaction tubes.
- 5. Cell-Saver tips (Biozyme Scientific).
- 6. Petri dishes (145 mm).
- 7. Folded filter paper.
- 8. Sterile filter units with  $0.22 \ \mu m$  pore size.
- 9. Incubator.
- 10. Rotary shaker.
- 11. Centrifuge with swing-out rotor; programmable settings should include the specification of acceleration/deceleration rates (*see* **Note 8**).
- 12. Benchtop centrifuge.
- Plasmid purification kit with prepacked gravity-flow anionexchange columns in maxi (500 µg) or mega (2.5 mg) scale;
   e.g. Plasmid Mega Kit (Qiagen), JETstar Plasmid Purification MIDI Kit (Genomed).
- 14. Fluid aspiration system.
- 15. Hemocytometer.
- 16. Vortexer.
- 17. FLUOstar Optima microplate reader (BMG Labtech) equipped with an on-board syringe injector (*see* **Note 1**).
- 18. 96-well microplates white LUMITRAC 200 (Greiner) (*see* Note 1).
- 19. 96-well microplates black FLUOTRAC 200 (Greiner) (see Note 1).
- 20. Nunc<sup>™</sup> MicroWell<sup>™</sup> 96-well microplates (Nunc) (*see* Note 1).
- 21. Equipment for maintenance of a cell suspension culture.
- 22. Nylon net filter with 70  $\mu$ m pore size (optional, *see* Note 10).

#### 2.3 Plasmids 1. Reporter constructs

Reporter constructs are based on the vector pBT10GUS [5] or the Gateway-compatible derivative pDISCO [10]. In both

cases the promoter (fragment) to analyze is inserted or recombined upstream of the *uidA* ORF:nos terminator cassette (*see* **Notes 2** and **3**).

In case of testing the activity of a transcriptional repressor, the construction of a weakly active synthetic promoter is needed (*see* **Note 4**).

2. Effector expression constructs

Effectors were expressed under the control of the strong constitutive CaMV 35S promoter (positions -417 to +8), inserted into classical pBT vectors [3] or the Gateway-compatible derivative pBTdest [6].

3. LUC standardization plasmid

The LUC standardization plasmid used in the cotransfection assays contains a *Photinus pyralis* luciferase (LUC) encoding open reading frame [11] under the control of the constitutive *Petroselinum crispum* UBI4-2 promoter [12] in pBT (*see* **Notes 5** and 6).

4. Filling plasmid

The promoter-deleted standardization plasmid pBT10- $\Delta$ -LUC, which in protoplasts leads to no detectable luciferase activity, is added to keep the total amount of the transfected plasmid DNA constant (25 µg).

#### 3 Methods

	This protocol uses <i>A. thaliana</i> At7 cells from a cell suspension cul- ture which is maintained at 26 °C in the dark on a rotary shaker at 105 rpm. Cells are subcultured once a week by transferring approx- imately 2.8 g of cells to 40 mL fresh MS-medium. For protoplast isolation, additional Erlenmeyer flasks with 40 mL MS-medium were inoculated 5 days prior to the harvesting of the cells.
3.1 Protoplast Preparation	1. For each subcultivated Erlenmeyer flask of At7 cells with 40 mL cell culture prepare 60 mL of cellulase-mazerozyme solution ( <i>see</i> <b>Note</b> 7).
	<ol> <li>Transfer each 5-day-old At7 cell suspension subculture to a 50 mL Falcon tube.</li> </ol>
	3. Centrifuge 5 min at $130 \times g$ with moderate acceleration (5/9) and deceleration (3/9) in a swing-out rotor ( <i>see</i> <b>Note 8</b> ).
	4. Discard the supernatant carefully.
	5. Detach cell pellet with caution by soft tapping.
	6. Add 50 mL 240 mM CaCl <sub>2</sub> and resuspend the cells by gentle inversion of the tube.
	7. Centrifuge 5 min at $130 \times g$ with moderate acceleration (5/9) and deceleration (3/9) in a swing-out rotor.



**Fig. 1** Schematic depiction of transient protoplast expression assay with co-transfected At7 protoplasts. The *gray box* at the *top* gives the plasmids (constructs) used in co-transfection experiments with the amounts and concentrations specified. The middle part depicts a transfected At7 protoplast with plasmids (*italics*) and the produced proteins (*normal letters*) and their interactions. The "filling" plasmid is not considered. The *brown box* at the *bottom* summarizes the calculation of normalized specific GUS' activity, as a measure for promoter activity. Abbreviations: *35S* cauliflower mosaic virus *35S* promoter, *CDS* coding sequence, *GUS*  $\beta$ -glucuronidase, *LUC* luciferase, *TF* transcription factor, *Ubi ubiquitin4-2* promoter

- 8. Discard the supernatant carefully.
- 9. For each harvested At7 cell suspension flask, prepare two 145 mm Petri dishes each with 10 mL cellulase-mazerozyme solution (*see* Note 7).
- 10. Little by little, add the residual 40 mL of the enzyme solution to the cell pellet and gently resuspend by inversion. Avoid cell clumping.
- 11. Add half of the cell suspension (about 20 mL) to each of the prepared Petri dishes.
- 12. Incubate overnight at 26 °C in the dark, shake at 20 rpm.
- 13. Intensify the shaking to 40 rpm for no longer than 20 min.
- From each Petri dish, carefully transfer the protoplasts into a 50 mL Falcon tube (*see* Note9).
- 15. Centrifuge 5 min at  $90 \times g$  with moderate acceleration (5/9) and deceleration (3/9) in a swing-out rotor.
- 16. Discard the supernatant carefully.
- 17. Detach the pellet with caution by soft tapping.
- Wash the protoplasts by little and little adding 25 mL 240 mM CaCl<sub>2</sub> and resuspend the protoplasts by gentle inversion of the tube (*see* Note 10).
- 19. Centrifuge 5 min at  $90 \times g$  with moderate acceleration (5/9) and deceleration (3/9) in a swing-out rotor.
- 20. Discard the supernatant carefully.
- 21. Detach the pellet with caution by soft tapping.
- 22. Little by little add 20 mL B-5 floating medium to each pellet and combine two resuspended pellets from 50 mL Falcon tubes in one tube (*see* Note11)
- 23. Centrifuge 5 min at  $130 \times g$  with maximal acceleration (9/9) and minimal deceleration (1/9) in a swing-out rotor.
- 24. Transfer the floating protoplasts (2–5 mL) with a Cell-Saver tip into a new 50 mL Falcon tube (*see* **Note12**).
- 25. Cautiously fill the Falcon tube with B-5 floating medium.
- 26. Centrifuge 5 min at  $130 \times g$  with maximal acceleration (9/9) and minimal deceleration (1/9) in a swing-out rotor.
- 27. Pool all floating protoplasts in a 13 mL centrifuge tube using a Cell-saver tip.
- 28. Assess the quality of the protoplast suspension (*see* Note 13).
- 29. Use the protoplasts immediately for transfection. 200  $\mu$ L of protoplasts (containing  $1-2 \times 10^6$  At7 protoplasts) are needed per co-transfection (*see* **Note 14**).

3.2 Preparing DNA for Co-transfection	<ol> <li>Plasmids to be used in protoplast co-transfection experiments are retransformed into the <i>dam</i> and <i>dcm</i> methylation-deficient <i>E. coli</i> strain K12 ER2925 (NEB) (<i>see</i> Note 15).</li> <li>Plasmid DNA is prepared from methylation-deficient <i>E. coli</i> strain using a plasmid purification kit with prepacked gravity- flow anion-exchange columns in maxi (500 µg) or mega (2.5 mg) scale according to manufacturer's instructions.</li> <li>The high concentrated <i>dam</i>- plasmid DNA should be stored at 4 °C (<i>see</i> Note 16).</li> <li>For co-transfection experiments, plasmid dilutions have to be made: reporter constructs, standardization constructs and "filling" constructs with 1 µg/µL, effector constructs with 0.1 µg/µL (Fig. 1).</li> <li>Combined plasmid DNA solutions for co-transfection should be prepared in advance to enable a smooth execution of the protocol. All combined DNA solutions should contain an equal amount of plasmid DNA (25 µg) in a volume of 50 µL (Fig. 1). The use of a positive control (35S::GUS reporter con- struct) and a negative control (TATA::GUS reporter construct, containing only the truncated -46 minimal promoter (TATA))</li> </ol>
3.3 Protoplast	<ul><li>is recommended in the experimental setup (<i>see</i> Note17).</li><li>1. Thaw PEG solution at room temperature (<i>see</i> Note18).</li></ul>
Co-transfection	2. For each co-transfection transfer 200 $\mu$ L of the protoplast suspension with a Cell-Saver tip into a 13 mL centrifuge tube.
	<ol> <li>Pipet the prepared DNA solutions (25 μg in 50 μL) onto the protoplasts (<i>see</i> Note 19).</li> </ol>
	4. Add 200 $\mu$ L PEG solution and mix thoroughly but gently by soft shaking and tapping of the tube rack ( <i>see</i> <b>Note 20</b> ).
	5. Incubate the protoplast–DNA–PEG mixture at room tempera- ture for 15 min ( <i>see</i> <b>Note 21</b> ).
	6. Stop the transfection reaction by stepwise adding 5 mL 275 mM $Ca(NO_3)_2$ (see Note 22).
	7. Centrifuge 5 min at $90 \times g$ with maximal acceleration (9/9) and moderate deceleration (5/9) in a swing-out rotor.
	8. Discard the supernatant carefully.
	9. Stepwise add 7 mL B5 solution (see Note 23).
	10. Incubate at 26 °C for approximately 20 h in the dark, keeping the tubes in an almost horizontal position ( <i>see</i> <b>Note 24</b> ).
3.4 Harvesting of Transfected	1. Prepare 50 mL Falcon tubes with 20 mL of cold 240 mM CaCl2 (4 °C) for each co-transfection.
Protoplasts	2. Add the protoplasts within the B5 solution to the tubes by decanting.

	3. Centrifuge for 10 min at $300 \times g$ and 4 °C in a swing-out rotor.
	4. Remove the supernatant with a fluid aspiration system down to ca. 1 mL.
	5. Resuspend the pellet in the remaining supernatant and transfer the protoplast suspension into a 1.5 mL reaction tubes using Cell-Saver tips.
	6. Centrifuge 30 s at $10,000 \times g$ and 4 °C.
	7. Remove the supernatant using the fluid aspiration system and instantly freeze the protoplast pellet in liquid nitrogen.
	8. Store the frozen protoplasts at –80 °C until use or thaw on ice for following protein extraction.
3.5 Extract Proteins	1. Thaw the protoplasts on ice.
from Protoplasts	2. Add 750 µL of protein extraction buffer.
	3. Resuspend the pellet by vortexing rigorously for a minimum of 30 s.
	4. Centrifuge for 10 min at full speed on 4 °C in a benchtop centrifuge.
	5. Keep the reaction tubes on ice; the supernatant is used for the protein and reporter gene assays ( <i>see</i> <b>Note 25</b> ).
3.6 Reporter Gene Assays	In this assay the luciferase activity is quantified as a measure for transfection efficiency ( <i>see</i> Note 25).
3.6.1 Luciferase Assay	<ol> <li>Pipet 10 μL of the protoplast protein extracts to the wells of a white LUMITRAC 96-well microplate.</li> </ol>
	2. Adjust the instrument settings to luminescence detection.
	3. Fill the syringe injector of the FLUOstar Optima microplate reader with 100 $\mu$ L luciferase substrate solution for each sample and start the luminescence measurement ( <i>see</i> Note 26).
	<ol> <li>Measure the produced light (relative light unit, RLU) during 10 s using the FLUOstar Optima microplate reader.</li> </ol>
	<ol> <li>Calculate the specific luciferase activity LUC<i>i</i> [RLU/s/μg] for each co-transfection by dividing the measured light [RLU/s/μL] by the protein concentration [μg/μL].</li> </ol>
	RLU
	$LUC_i = \frac{\overline{s \times \mu L}}{\mu g \text{ protein}}$

μL

- 3.6.2 GUS Assay Fluorometric analysis allows quantification of GUS activity. In the presence of GUS, MUG is hydrolyzed to the fluorescent product 4-methylumbelliferone (MU). After the reaction, total fluorescence is measured and product concentration is calculated based on a MU standardization curve.
  - 1. Pipet 100  $\mu$ L of the protein extracts into the wells of a black FLUOTRAC 200 96-well microplate.
  - 2. Add 100  $\mu L$  of 4-MUG substrate solution to each protein extract.
  - 3. Pipet the MU dilution series to the microplate as well. This series is used to generate a MU standardization curve.
  - 4. Set excitation to 365 nm and read the sample emission at 455 nm after 20, 40, and 60 min at 37 °C in the FLUOstar Optima microplate reader.
  - 5. Determine the average change in measured MU fluorescence  $(\Delta E_{455})$  from 20 to 40 min, and from 40 to 60 min (see Note 27).
  - 6. Calculate the line of best fit for the MU fluorescence in the MU dilution series (MU standardization curve). Determine the slope (*m*) of the MU standardization curve.
  - 7. The specific β-glucuronidase activity GUS*i* [pmol/min/mg] is determined by the formula:

$$GUS_{i} = \frac{\Delta E_{455} \times 1000 \frac{\mu g}{mg} \times \frac{200 \mu L}{20 \mu L}}{20 \min \times \frac{m}{pmol} \times \mu g \text{ protein}}$$

The GUS activity measurement uses 20  $\mu$ L of a 200  $\mu$ L sample (consisting of 100  $\mu$ L protein extract and 100  $\mu$ L 4-MUG substrate solution).

The amount of protein  $[\mu g]$  in 100  $\mu L$  protein extract is given.

To determine the protein concentration in the protein extract samples, a Bradford assay [13] is performed.

- 1. Pipet 10 μL of the protein extracts to the wells of a Nunc<sup>™</sup> MicroWell<sup>™</sup> 96-well microplate.
- 2. Pipet 10  $\mu$ L of the BSA dilution series to the microplate as well. This series is used to generate a protein standardization curve.
- 3. Mix with 200  $\mu$ L protein assay dye reagent.

3.7 Protein Concentration Measurement

- 4. Incubate for 5 min at 37 °C.
- 5. Measure OD<sub>595</sub> in the FLUOstar Optima microplate reader.
- 6. Calculate the protein concentrations in the extracts with help of the BSA dilution series.
- 7. Determine the amount of protein  $[\mu g]$  in 10  $\mu$ L protein extract and calculate the protein concentration  $[\mu g/\mu L]$ .

3.8 Calculation of the Normalized Specific GUS' Activity (See Note 28)

- 1. We normally repeat each co-transfection experiment (with the same combination of plasmids) six times, with six independent (*i*) co-transfections with three different protoplast preparations, giving an "experimental block". A "whole experiment", including controls and all related experiments to answer a biological question, consists of several experimental blocks.
- 2. Calculate the average of all specific LUC*i* values (LUC*M*) from a whole experiment.

$$LUC_M = \frac{1}{n} \times \sum LUC_i$$

*n*: sum of all co-transfections in a whole experiment

3. For standardization, a specific correction factor *Fi* for each individual co-transfection experiment is determined by dividing LUC*M* by the specific LUC*i* value.

$$F_i = \frac{\text{LUC}_M}{\text{LUC}_i}$$

4. The standardized, corrected GUS activity (GUS*ki*) is obtained by multiplying the specific correction factor *Fi* with the specific GUS activity GUS*i*.

$$GUS_{ki} = F_i \times GUS_i$$

5. The average of specific GUS*ki* values of an experimental block of six co-transfections is calculated as standardized GUS activity (GUS').

$$GUS' = \frac{1}{6} \times \sum GUS_{ki}$$

6. The standard deviation of GUS' (SD(GUS')) is determined by the formula:

$$\mathrm{SD}(\mathrm{GUS}') = \frac{1}{\sqrt{6(6-1)}} \times \sqrt{\sum (\mathrm{GUS}_{ki} - \mathrm{GUS}')^2}$$

#### 4 Notes

- 1. It is also possible to use alternative reaction containers and instruments for the protein, LUC, and GUS measurements.
- The transfection rate can be variable and depends on the plasmids used [5]. Generally, small vector sizes achieve higher transformation rates. The pBT vectors have a size-minimized backbone of 1989 bp containing colE1-ori and amp<sup>r</sup> [3].
- 3. *cis*-acting elements often contain palindromic sequences. This has been interpreted as a reflection of the fact that DNAbinding proteins are in many cases active as dimers or tetramers, binding symmetrically to DNA matching the symmetry of the binding site [14]. Otherwise, several asymmetric *cis*elements are known which are often recognized by heterodimeric *trans*-acting factors. Aiming to test the interaction of synthetic *cis*-elements and DNA binding proteins, we found it helpful to insert such binding elements in both orientations single copy, as a dimer, and as a tetramer fused to the GUS coding sequence in the pBT10GUS vector.
- 4. A promoter with appreciable activity in At7 protoplasts in the absence of added effectors has been reported. This reporter construct contains the region between -90 to +8 of the CaMV 35S promoter (-90 CaMV 35S promoter), including an activation sequence factor 1 binding site (positions -83 to -63) [15], fused to *uidA* encoding GUS. Binding domains (*cis*-elements) to analyze can be cloned immediately upstream of the -90 CaMV 35S promoter in pBT10GUS [4].
- 5. Addition of the LUC standardization plasmid in the co-transfection is used to determine specific LUC activity in a given sample to estimate the transfection rate (efficiency of transfection).
- 6. The *Photinus pyralis* luciferase (LUC) encoding ORF contains three silent point mutations which remove XbaI, EcoRI, and ClaI sites [16].
- 7. We observed strong differences between different lots of cellulase in terms of success in protoplasting. Each lot has to be tested and the amount of cellulase in the cellulase-mazerozyme solution has to be adjusted for successful protoplasting. Mazerozym is a multi-component enzyme mixture containing activities of pectinase,  $\alpha$  amylase and hemicellulase; cellulase proteins hydrolyze 1,4- $\beta$ -D-glucosidic bonds.
- 8. We are using a Multifuge 1s (Heraeus) centrifuge with a TTH400 swing-out rotor (Heraeus) or a Multifuge 3s-r (Heraeus) centrifuge with a TTH750 swing-out rotor (Heraeus). These centrifuges have the option to specify acceleration and deceleration in ramps from 1 to 9. These parameters have been optimized to protect the delicate living samples.

- 9. When harvesting the protoplasts from the Petri dishes, transfer the cell suspension into the Falcon tube by placing the brim of the Petri dish centrally over the tube and then decant carefully. Work above the inverted lid of the Petri dish to be able to recover eventually spilled protoplast suspension.
- 10. When adding solutions to the protoplasts, never just pour the liquid on top of the protoplasts, rather hold the tube at a flat angle and let the liquid slowly run along the side of the tube. Slow rotation of the tube helps dissolving solid pellets.
- 11. By washing with B5 floating medium, the living protoplasts are separated from the debris of broken cells. The high sugar concentration in the B5 floating medium causes floating of intact protoplasts.
- 12. When pipetting living protoplasts, always use pipetting equipment which reduces the shearing force, such as Cell-Saver tips or pipettes with a wide tip orifice.
- 13. When the floating protoplast suspension contains a high amount of cell clusters (caused by inefficient enzyme treatment for cell wall removal), it is possible to filter the protoplasts through a nylon net filter with 70  $\mu$ m pore size. Although this filtering step drastically reduces the number of cells, the obtained protoplast suspension contains only the desired protoplasts.
- 14. Protoplasts number is determined using a hemocytometer.
- 15. The use of *dam* plasmid DNA results in significantly reduced background activity, as shown for parsley protoplasts [17].
- 16. In our hands, high-concentrated (>1  $\mu$ g/ $\mu$ L) *dam* plasmid DNA in TE (10 mM Tris–HCl pH 8, 1 mM EDTA) stored at 4 °C is stable for more than 10 years, and can be used in co-transfection experiments without known limitations.
- 17. We recommend to generate and use a detailed pipetting plan.
- 18. Thawing of PEG solution at room temperature could take a few hours.
- 19. In order to ensure a simultaneous start of co-transfections for all experiments, do not mix the DNA and protoplasts at this point. If you are performing a small-scale experiment this is not as critical as if you are handling 40 co-transfection reactions at the same time.
- 20. When using a dispenser to add the PEG solution, it is recommended to hold the tubes or the rack with the tubes at a flat angel and pipet to the side of the tube.
- 21. Avoid to move the tubes during the incubation time.
- 22. We found that splitting the pipetting of  $Ca(NO_3)_2$  into two steps worked well to achieve a stopping of all reactions at

around the same time. First pipet 1 mL to all the tubes and then add the remaining 4 mL.

- 23. Split the pipetting of the 7 mL in at least two steps and first pipet 1 mL to all tubes. If you add the whole volume of B5 solution at once, it might result in clumping of the protoplasts.
- 24. Almost horizontal incubation avoids contact of the liquid to the cap of the centrifugation tube, which might result in protoplasts stuck to the cap.
- 25. The luciferase assay should be performed immediately after protein extraction as the luciferase is degraded in the extract. The recommended order in which the following measurements have to be performed is (1)luciferase assay, (2)GUS assay, and (3)protein concentration measurement.
- 26. The luminescence measurement should immediately start after the addition of luciferase substrate solution. We implemented a program on the FLUOstar Optima microplate reader that adds 100  $\mu$ L of luciferase substrate solution to the samples immediately before the luminescence is measured.
- 27. Only assays with linear increase of the  $E_{455}$  values are taken into account.
- 28. We recommend the generation and use of an *Excel* sheet for the calculations.

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