Characterisation of a *PR-1*::Luciferase Transgenic Line Deployed to Uncover Novel Defence-Related Arabidopsis Mutants by Luciferase Imaging

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Abstract

In order to identify components of the defence signalling network that may contribute to the establishment of disease resistance, we generated a novel *PR-1::Luciferase* transgenic line which was deployed in an imaging based screen to uncover novel defence-related mutants. Approximately, 5 000 ethylmethane sulfonate (EMS) lines and 30 000 activation tagged lines were generated and screened for enhanced LUC activity via ultra low light imaging.

Keywords: signal transduction; luciferase imaging; systemic acquired resistance; Arabidopsis mutants

INTRODUCTION

Plants have evolved an extensive array of sophisticated defence mechanisms which are employed to resist the establishment of disease (GLAZEBROOK et al. 1997). Pre-formed physical and chemical barriers constitute the first line of defence (OSBOURN 1996), upon which are superimposed a battery of inducible defence responses, including a rapidly activated oxidative burst (GRANT & LOAKE 2000), the cross-linking of cell wall structural components (WALLACE & FRY 1999) and the accumulation of a variety of anti-microbial proteins (BOL et al. 1990). The expeditious engagement of these responses is dependent upon successful pathogen recognition, which is thought to occur following the interaction of a specific pathogen avirulence (avr) gene product with a corresponding plant resistance (R) gene product (DANGL 1995). A near ubiquitous feature of successful pathogen recognition is the visible development of hypersensitive cell death (HR) surrounding the area of attempted pathogen ingress (GREENBERG 1997). Following HR development, is the establishment of immunity throughout the plant to a broad spectrum of ordinarily virulent pathogens. This phenomenon has been termed systemic acquired resistance (SAR) (RYALS et al. 1996). The establishment of SAR correlates with a rise in peroxidase activity, increased lignin deposition and the expression of so-called SAR genes, which encode pathogenesis-related (PR) proteins.

MATERIAL AND METHODS

The PR-1::LUC::OCS chimeric gene was constructed as described in Murray et~al.~(2002). The generated pART27 binary vector was then transformed into Agrobacterium~tumefaciens strain GV3101. Arabidopsis ecotype Col-0 was then transformed by vacuum infiltration and transgenic plants were identified following growth in MS medium containing kanamycin. Pseudomonas~syringae~pv~tomato~DC3000~(PstDC3000) was grown as described in Murray et~al.~(2002). Five week old soil-grown plants were infected with a $Pst~DC3000~suspension~(OD_{600}=0.002)~in~10~mM$ MgCl₂ by completely infiltrating the abaxial side of the leaf with a 1 ml syringe.

RESULT

In order to accurately determine in real time, in living plant tissue, the temporal and spatial expression programme established by the *PR-1* gene during the

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establishment of R gene-mediated resistance and SAR, we generated a transgenic Arabidopsis line containing a chimeric PR-1::LUC gene fusion. The PR-1 promoter was transcriptionally fused to the firefly luciferase (LUC) reporter gene (Figure 1A), and the resulting construct was transformed into Arabidopsis accession Col-0. To confirm the utility of a typical transgenic line, the profile of PR-1::LUC gene expression was compared to that of the endogenous PR-1 gene following inoculation of Pseudomonas syringae pv. tomato (Pst) DC3000 expressing the avrB gene (Figure 1B). From 12 hours post inoculation, expression of the PR-1::LUC gene steadily increased, reaching a maximum value between 36 and 48 hours post inoculation, after which time *PR-1*::*LUC* expression slowly decreased. The expression of the endogenous PR-1 gene was found to be congruent with expression of the PR1: :LUC transgene. We also determined the temporal profile of LUC activity following inoculation with Pst DC3000 (avrB) (Figure 1C). This also exhibited similar kinetics to that of the endogenous PR-1 gene, with maximum LUC activity recorded at 36 h post

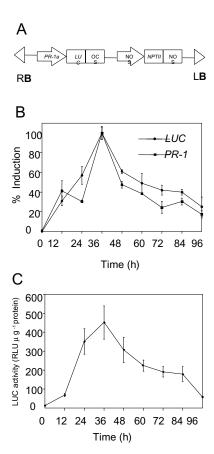


Figure 1. LUC activity following inoculation with *Pseudomonas syringae* pv. *tomato* DC3000

inoculation. These experiments therefore confirmed that the *PR-1*::*LUC* transgene functioned as a robust reporter for the engagement of endogenous *PR-1* gene expression.

DISCUSSION

We have generated an *Arabidopsis* line containing a chimeric *PR-1::LUC* transgene that robustly reports *PR-1* gene expression during the establishment of *R* gene-mediated disease resistance and SAR. The utility of the novel *PR-1::LUC* transgenic line was exploited in a screen designed to uncover EMS *Arabidopsis* mutants that caused miss-expression of the *PR-1* gene (Murray *et al.* 2002). To uncover potentially redundant and/or essential genes integral to the establishment of SAR in *Arabidopsis*, we generated a large collection of activation tagged lines and screened via ultra low light imaging.

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