A Universal Role for Inositol 1,4,5-Trisphosphate-Mediated Signaling in Plant Gravitropism


Department of Botany, North Carolina State University, Raleigh, North Carolina 27695–7612 (I.Y.P., C.-Y.H., W.F.B.); and Department of Biology, Wake Forest University, Winston-Salem, North Carolina 27109–7325 (S.B., G.K.M.)

Inositol 1,4,5-trisphosphate (InsP$_3$) has been implicated in the early signaling events of plants linking gravity sensing to the initiation of the gravitropic response. However, at present, the contribution of the phosphoinositide signaling pathway in plant gravitropism is not well understood. To delineate the role of InsP$_3$ in plant gravitropism, we generated Arabidopsis (Arabidopsis thaliana) plants constitutively expressing the human type I inositol polyphosphate 5-phosphatase (InsP$_5$-ptase), an enzyme that specifically hydrolyzes InsP$_5$. The transgenic plants show no significant differences in growth and life cycle compared to wild-type plants, although basal InsP$_3$ levels are reduced by greater than 90% compared to wild-type plants. With gravistimulation, InsP$_3$ levels in inflorescence stems of transgenic plants show no detectable change, whereas in wild-type plant inflorescences, InsP$_3$ levels increase approximately 3-fold within the first 5 to 15 min of gravistimulation, preceding visible bending. Furthermore, gravitropic bending of the roots, hypocotyls, and inflorescence stems of the InsP$_5$-ptase transgenic plants is reduced by approximately 30% compared with the wild type. Additionally, the cold memory response of the transgenic plants is attenuated, indicating that InsP$_3$ contributes to gravitropism in the cold. The transgenic roots were shown to have altered calcium sensitivity in controlling gravitropic response, a reduction in basipetal indole-3-acetic acid transport, and a delay in the asymmetric auxin-induced β-glucuronidase expression with gravistimulation as compared to the controls. The compromised gravitropic response in all the major axes of growth in the transgenic Arabidopsis plants reveals a universal role for InsP$_3$ in the gravity signal transduction cascade of plants.

Both gravity and light direct plant growth so that shoots grow upward for efficient photosynthesis and gas exchange and roots grow downward into the soil for anchorage and for water and mineral uptake (Hangarter, 1997). When the orientation of a plant is altered with respect to the gravity vector (i.e. when a plant is blown over by wind or rain), the shoots and roots exhibit differential growth resulting in upward and downward curvature, respectively, which returns the plant to a vertical orientation.

Gravitropism is a universal plant response. All major axes of plant growth, including roots, hypocotyls, coleoptiles, and inflorescence stems, are capable of a gravitropic response, despite differences in their basic structural organization (Tasaka et al., 1999). The presence of mutants defective in the gravity response of either the root or shoot suggests that the mechanisms of gravitropism in roots and shoots have both common and separate steps (Lomax, 1997; Tasaka et al., 1999). Nevertheless, the overall process of gravitropism is well conserved and consists of three major phases—gravity sensing, information transfer from sensing to responding cells (signal transduction and amplification), and the growth response (for review, see Perbal and Driss-Ecole, 2003; Morita and Tasaka, 2004).

One of the primary events in gravity sensing in higher plants involves the displacement or settling of dense starch-containing amyloplasts, which can occur in seconds to minutes depending on the plant tissue (Sack, 1991, 1997). Amyloplasts are found in specialized cells, such as the starch parenchyma surrounding the vascular tissue in plant shoots and grass pulvini, and in the columella of the root cap (Sack, 1991; Chen et al., 1999; Rosen et al., 1999; Ferrin et al., 2005). Upon reorientation of a plant shoot or root, the settling of amyloplasts, or the pressure exerted by protoplast settlement (Staves, 1997), is thought to trigger intra- and intercellular signaling, initiating downstream metabolic changes, including a lateral redistribution of auxin, leading to an asymmetric growth response (Lomax et al., 1995; Muday and DeLong, 2001).

Over the past few years, cumulative evidence has supported the role of amyloplasts in gravity sensing (Kiss, 2000; Blancaflor and Masson, 2003), although there is some debate about how the displacement of amyloplasts might trigger a signaling cascade. Some models propose that the amyloplasts are tethered by the actin cytoskeleton, and even small local perturbations...
in amyloplast movement could disrupt the actin network. This disruption may be transmitted to the plasma membrane (PM) and other endomembranes activating mechanosensitive channels (Volkman and Baluska, 1999; Yoder et al., 2001). The role of the actin cytoskeleton in gravitropism appears to be more complex, as shown by recent reports by Yamamoto and Kiss (2002) and Hou et al. (2003), and may act to down-regulate gravisignaling (Hou et al., 2004).

Many signaling molecules and second messengers, such as Ca\(^{2+}\) (Pieth and Trewavas, 2002), inositol 1,4,5-trisphosphate (InsP\(_3\); Perera et al., 1999, 2001), the Cholodny-Went hypothesis, lateral gradients of the plant auxin indole-3-acetic acid (IAA) have been suggested to induce this differential growth. More recently, asymmetric distribution of either endogenous free IAA or exogenously applied radiolabeled IAA across gravity-stimulated roots and shoots has been demonstrated in several plant systems, including gravity-stimulated coleoptiles (Parker and Briggs, 1990; Philippar et al., 1999), pulvini tissues (Long et al., 2002), and roots (Young et al., 1990), and has been shown to precede gravitropic bending (Parker and Briggs, 1990). Gradients of IAA have also been indirectly visualized using auxin-responsive promoters to drive the expression of reporter genes, such as \(\beta\)-glucuronidase (GUS) in shoots (Li et al., 1991) and in roots (Rashotte et al., 2001; Buer and Muday, 2004; Hou et al., 2004). A role for auxin transport in the gravitropic response is further supported by the fact that several auxin transport mutants show altered gravitropic responses and the application of IAA transport inhibitors to wild-type roots can abolish the gravitropic response (Muday, 2001; Blancaflor and Masson, 2003).

In previous work, we characterized the involvement of phosphoinositide (PI)-based signaling in the gravitropic response of cereal grass stems. Biphasic changes in InsP\(_3\) were detected with gravistimulation of maize and oat (Avena sativa) pulvini (Perera et al., 1999, 2001). These included rapid and transient fluctuations in InsP\(_3\) within seconds of gravistimulation in both upper and lower pulvini halves, followed by a long-term sustained increase in InsP\(_3\) only in lower pulvini halves. The second increase in InsP\(_3\) correlated with the bending response (Perera et al., 1999). Blocking the differential InsP\(_3\) gradient with a chemical inhibitor of phospholipase \(C\) (PLC) activity (U73122) attenuated the gravitropic response. Furthermore, the biphasic changes in InsP\(_3\) only occurred in the sensing and responding tissue (pulvinus) and were not detected in the non-elongating internodal regions of the stem (Perera et al., 2001). These results indicated that InsP\(_3\) plays a critical role in gravity signaling in cereal grass pulvini. However, it was not known whether InsP\(_3\) played a signaling role in both monocot and dicot plant gravitropism or whether the involvement of InsP\(_3\) was unique to the gravitropic response of the pulvinus system.

To test the hypothesis that InsP\(_3\) is a universal component involved in establishing the differential growth response in plants, we undertook a molecular approach to alter InsP\(_3\) signaling in Arabidopsis. We have generated transgenic Arabidopsis plants expressing the human type I inositol polyphosphate 5-phosphatase (InsP\(_5\)-ptase), an enzyme that specifically hydrolyzes the soluble inositol phosphates InsP\(_3\) and InsP\(_4\) (Laixinarayan et al., 1993). We chose a heterologous enzyme for several reasons. First, the animal type I InsP\(_5\)-ptase is well characterized and specifically hydrolyzes InsP\(_3\) and not the inositol phospholipids. Arabidopsis contains a family of related inositol 5-phosphatases known as At5PTases (Berdy et al., 2001), which have been shown to have roles in diverse aspects of plant growth and development, including vascular patterning in leaves and cotyledons (Carland and Nelson, 2004; Lin et al., 2005), secondary cell wall and fiber cell development (Zhong et al., 2004), and abscisic acid (ABA)-regulated processes (Sanchez and Chua, 2001; Burnette et al., 2003). However, there does not appear to be a homolog of the animal type I enzyme based on biochemical activity and sequence similarity. Several of the many plant inositol 5-phosphatases characterized to date can hydrolyze both lipid substrates and soluble inositol phosphates and are not specific for InsP\(_3\) hydrolysis (Berdy et al., 2001; Ercetin et al., 2004). Finally, the animal type I InsP\(_5\)-ptase is associated with the PM and therefore should preferentially dampen InsP\(_3\) signals generated at the PM.

We tested the feasibility of this approach by first generating transgenic tobacco (Nicotiana tabacum) cells constitutively expressing the human type I InsP\(_5\)-ptase (Perera et al., 2002). The transgenic tobacco cells had normal morphology and growth compared to wild type; however, basal InsP\(_3\) and phosphatidylinositol 4,5-bisphosphate (PtdInsP\(_2\)) levels were greatly reduced in the transgenic cells, resulting from the increased turnover of PtdInsP\(_3\) and an increased flux through the PI pathway.

In this article, we characterize the transgenic Arabidopsis lines expressing InsP\(_5\)-ptase. The plants show
no morphological differences compared to wild-type plants under normal growth conditions. Significantly, although vertical growth rates were comparable between wild-type and transgenic seedlings, the reorientation of transgenic roots and hypocotyls in response to gravistimulation was reduced by 30% compared to wild-type seedlings. Furthermore, the gravitropic response of inflorescence stems of transgenic plants was also attenuated compared to wild type. InsP$_3$ levels were found to increase in wild-type inflorescence stems upon gravistimulation, preceding visible bending, consistent with our previous results on cereal grass pulvini. No increases in InsP$_3$ with gravistimulation were detectable in the transgenic plants. In the transgenic roots, basipetal auxin transport was reduced and the development of the lateral auxin asymmetry on the lower side of gravistimulated roots was delayed. Our results indicate that dampening of the InsP$_3$ signal attenuates the gravitropic response and suggest that InsP$_3$-mediated signaling is a necessary component for the full gravitropic response in a dicot system. Additionally, since the PI pathway is implicated in plant responses to many different stresses (Stevenson et al., 2000; Meijer and Munnik, 2003), we anticipate that the transgenic plants will be a useful model system to evaluate the involvement of InsP$_3$-mediated signaling in plant responses to many different abiotic and biotic stresses.

RESULTS

Generation of Transgenic Lines Expressing InsP 5-ptase

Arabidopsis (Columbia-0) was transformed by Agrobacterium-mediated transformation using vacuum infiltration with the construct shown in Figure 1A. The coding region of the human type I InsP 5-ptase cDNA with a 6-His tag at the N terminus was inserted downstream of the cauliflower mosaic virus 35S promoter for constitutive expression. Four independent homozygous transformed lines were isolated by screening on selective media. The stable integration of a single copy of the transgene was verified by a Southern blot (data not shown). All growth and bending experiments were carried out using T$_4$ generation plants.

As seen in Figure 1, B and C, the transgenic plants grow normally and do not exhibit any significant morphological differences compared with the wild-type plants. The growth of the plants was monitored at various developmental stages from seed to seed as described by Boyes et al. (2001). Several vegetative and reproductive features, along with the timing of growth stages and plant yield, were compared between the wild type, vector control, and three independent transgenic lines (Fig. 1C; Table I). No significant differences were observed in any of the parameters measured, indicating the absence of pleiotropic effects of the transgene insertion.

The expression of the transgene was examined by immunoblot of total cellular protein extracts from 2-week-old transgenic and control plants. The InsP 5-ptase gene product is produced in all the transgenic lines tested (Fig. 2A). In addition, the transgene is expressed in all tissues tested, including roots, hypocotyls, mature leaves, and inflorescence stems of the transgenic plants (data not shown). Further fractionation of the cellular proteins by two-phase partitioning into soluble, lower phase, and PM-enriched fractions shows that the InsP 5-ptase protein is primarily associated with the PM fraction (Fig. 2B). We also monitored InsP 5-ptase enzyme activity in the transgenic
After 24 h, wild-type and vector control plants show a 2-fold increase in InsP3 hydrolysis activity compared to the PM fraction of wild-type plants (432.5 pmol min⁻¹). The specific activity in the PM-enriched fraction of transgenic plants is approximately 75-fold higher as compared to the PM activity in the PM-enriched fraction of transgenic plants described previously (Perera et al., 2002). The specific activity was quantified by measuring InsP3 hydrolysis using the soluble and PM fractions as described previously (Perera et al., 2002). The specific activity in the PM-enriched fraction of transgenic plants is approximately 75-fold higher as compared to the PM fraction of wild-type plants (432.5 pmol min⁻¹ protein for transgenic PM and 5.65 pmol min⁻¹ protein for wild-type PM). In addition, the PM fraction of the transgenic plants has approximately 5-fold higher specific activity than the soluble fractions from the same plants (data not shown). The localization of InsP5-ptase protein and enzyme activity are consistent with the PM localization of the InsP5-5ptase protein in animal cells (De Smedt et al., 1996).

Consistent with stable constitutive gene expression and elevated enzyme activity, basal InsP3 levels were found to be drastically reduced in the transgenic lines as compared with the wild type and vector control (Table III). InsP3 levels in roots and hypocotyls of 7-d-old seedlings were measured using the receptor-binding assay (Perera et al., 2002). In wild-type seedlings, InsP3 content is typically in the range of 400 to 600 pmol g⁻¹ fresh weight and is reduced by approximately 95% in the transgenic lines to 20 to 30 pmol g⁻¹ fresh weight.

### Table 1. Growth-stage measurements

<table>
<thead>
<tr>
<th>Plant Line</th>
<th>Total No. of Rosette Leaves</th>
<th>Length of Longest Leaf</th>
<th>Area of Rosette</th>
<th>Length of Primary Stalk</th>
<th>Seed Production</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wild type</td>
<td>8.8 ± 1.8</td>
<td>33.7 ± 0.3</td>
<td>32.7 ± 3.8</td>
<td>41.3 ± 0.7</td>
<td>43.8 ± 1.8</td>
</tr>
<tr>
<td>C2</td>
<td>10.2 ± 2.0</td>
<td>33.5 ± 1.1</td>
<td>32.7 ± 2.5</td>
<td>39.4 ± 2.4</td>
<td>34.6 ± 6.7</td>
</tr>
<tr>
<td>2-6</td>
<td>10.1 ± 1.6</td>
<td>34.6 ± 1.9</td>
<td>33.5 ± 5.0</td>
<td>35.6 ± 1.2</td>
<td>41.7 ± 4.3</td>
</tr>
<tr>
<td>2-8</td>
<td>8.9 ± 1.2</td>
<td>31.8 ± 1.0</td>
<td>28.6 ± 2.0</td>
<td>36.2 ± 3.6</td>
<td>36.6 ± 10.3</td>
</tr>
<tr>
<td>2-12</td>
<td>9.5 ± 1.1</td>
<td>31.7 ± 0.1</td>
<td>28.6 ± 3.5</td>
<td>37.3 ± 2.4</td>
<td>36.3 ± 9.2</td>
</tr>
</tbody>
</table>

The values are the average of two independent experiments ±sd.

Growth and Gravitropic Responses of Roots and Hypocotyls

To determine whether the transgenic plants with severely reduced InsP3 levels would show an altered response to gravity, the growth and gravitropic responses were monitored in roots of 4-d-old seedlings. Seedlings were grown vertically on agar plates in the dark and vertical elongation growth was monitored over a 24-h period. There were no significant differences in vertical growth rates between wild-type, vector control, and transgenic lines expressing InsP5-5ptase (Table III). In addition, staining with I2 has shown that starch accumulation in amyloplasts of root cap columella cells is normal in both wild-type and transgenic roots (data not shown). To monitor the gravitropic response, plates were turned by 90° in the dark and images were recorded over a period of 24 h. Bending was first visible after approximately 2 h and, by 6 h of gravistimulation, wild-type and vector control plants exhibit approximately 30° to 35° curvature (Fig. 3A). After 24 h, wild-type and vector control plants show approximately 70° curvature. In contrast, the transgenic lines consistently have a reduced degree of curvature at both 6 and 24 h of gravistimulation. Compared to wild-type roots, the gravitropic response of the transgenic roots is reduced by approximately 30%.

To examine the early stages of the root gravitropic response in more detail, we analyzed curvature using Multi-ADAPT software developed by Ishikawa and Evans (1997). This program provides high spatial and temporal resolution of the gravitropic bending response of a single root by measuring both root tip angle and elongation rates on the upper and lower flanks of the root. Upon gravistimulation, wild-type roots exhibit a lag period of approximately 120 min prior to initiation of curvature (Table IV). This is in good agreement with values previously published (Buer and Muday, 2004). The transgenic roots appear to have a much greater variability in initiation of bending (note the sd for this response is twice as large as for the wild type). The transgenic roots also have a slightly longer lag compared with the wild type, although this difference is not statistically significant due to the variability in the response of the transgenic roots. The most pronounced difference between the wild-type and transgenic roots is in the rate of curvature after bending is initiated.

![Figure 2](https://www.plantphysiol.org)
Transgenic roots have an approximately 45% reduction in rate compared to wild type, which would help explain the reduced gravitropic response observed in the roots.

We also evaluated growth and the gravitropic response of hypocotyls of young seedlings. Arabidopsis seedlings were grown vertically on agar plates in the dark and hypocotyl elongation was measured over a 48-h period. No significant differences were observed in vertical growth of hypocotyls of transgenic and wild-type plants (Table III). In contrast, the gravitropic bending response was reduced in hypocotyls of 4-d-old transgenic seedlings compared to wild type (Fig. 3B). Three independent transgenic lines (2-6, 2-8, and 2-12) showed an approximately 30% reduction in angle of curvature compared with the wild type and vector control lines. In all plants, hypocotyl reorientation in response to gravistimulation was slower than the root response. After 48 h of gravity stimulation, roots had mostly returned to a vertical orientation. Hypocotyls reached a final bending angle of approximately 45° after 48 h.

Although changes in InsP$_3$ in plants are associated with various stimuli and stresses, the downstream consequences of the InsP$_3$ changes are virtually unknown. InsP$_3$ is known to trigger the release of Ca$^{2+}$ from intracellular stores such as the vacuole and endoplasmic reticulum (for review, see Sanders et al., 1999).

Furthermore, the generation of InsP$_3$ may require Ca$^{2+}$ because all known plant PLC enzymes are regulated by Ca$^{2+}$ (Hunt et al., 2004). Therefore, there seems to be an interdependence and close connection between InsP$_3$ and Ca$^{2+}$. To investigate the relationship between Ca$^{2+}$ and InsP$_3$, further, we analyzed the growth and gravitropic responses of roots and hypocotyls under different Ca$^{2+}$ concentrations. Murashige and Skoog (MS) medium was made by omitting the Ca$^{2+}$, which will be referred to as “no added Ca$^{2+}$.” This medium may contain trace amounts of Ca$^{2+}$ as contaminants in the medium salts and the agar. The highest Ca$^{2+}$ concentration tested was 10 mM; standard MS medium contains 3 mM Ca$^{2+}$. The vertical growth of wild-type and transgenic seedlings was first monitored on plates containing the medium described above. As seen with the seedlings grown on regular MS medium, there were no differences in vertical root elongation between wild-type and transgenic seedlings over a 3-d period on both the no added Ca$^{2+}$ and 10 mM Ca$^{2+}$ (Fig. 4A). All of the roots (wild type and transgenic) showed decreased growth on the medium with no added Ca$^{2+}$ (similar growth over a 3-d period as was seen in 24 h on regular MS medium; Table III). All roots (wild type and transgenic) showed increased growth on the 10 mM Ca$^{2+}$ medium compared to the no added Ca$^{2+}$ medium, suggesting that, in the range of 3 to 10 mM, Ca$^{2+}$ promoted root growth. We then monitored root bending (Fig. 4, B and C). There was a statistically significant difference in root bending of all the lines examined at the two different Ca$^{2+}$ concentrations (10 mM or no added Ca$^{2+}$). With both treatments, the transgenic roots exhibited less bending than the wild type and vector control, consistent with our previous results on regular MS medium. However, when the extent of bending was compared between these two different Ca$^{2+}$ concentrations, the results were quite intriguing. While root bending was enhanced at 10 mM Ca$^{2+}$ for the wild type and vector control, the transgenic roots showed the opposite effect and exhibited less bending at 10 mM Ca$^{2+}$ than on the no added Ca$^{2+}$ medium. This difference was noticeable at both the 6- and 24-h time points (Fig. 4, B and C). After 48 h, however, the transgenic roots on the 10 mM Ca$^{2+}$ medium were not significantly different from the roots grown on medium with no added Ca$^{2+}$. This suggests that the high Ca$^{2+}$ decreases the reorientation response of the transgenic roots most dramatically within the first few hours.

Similar results were obtained with hypocotyl growth and gravitropism on altered Ca$^{2+}$ medium. Hypocotyls of 4-d-old wild-type and transgenic seedlings grew similarly on both no added Ca$^{2+}$ and 10 mM Ca$^{2+}$ with greater elongation at the 10 mM Ca$^{2+}$. When hypocotyl bending was monitored, wild-type and vector control

<table>
<thead>
<tr>
<th>Plant Line</th>
<th>InsP$_3$ Content$^a$</th>
<th>Root</th>
<th>% wild type</th>
<th>Hypocotyl</th>
<th>% wild type</th>
</tr>
</thead>
<tbody>
<tr>
<td>Wild type</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
<td>100</td>
</tr>
<tr>
<td>C2</td>
<td>107</td>
<td>106</td>
<td>3</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>2-6</td>
<td>3</td>
<td>3</td>
<td>2</td>
<td>2</td>
<td></td>
</tr>
<tr>
<td>2-8</td>
<td>2</td>
<td>2</td>
<td>5</td>
<td>5</td>
<td></td>
</tr>
<tr>
<td>2-12</td>
<td>9</td>
<td>5</td>
<td>6</td>
<td>6</td>
<td></td>
</tr>
</tbody>
</table>

$^a$The data presented are the average from two independent experiments.

<table>
<thead>
<tr>
<th>Plant Line</th>
<th>Length$^b$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Root</td>
<td>mm</td>
</tr>
<tr>
<td>Hypocotyl</td>
<td>mm</td>
</tr>
<tr>
<td>Wild type</td>
<td>3.5 ± 0.4</td>
</tr>
<tr>
<td>C2</td>
<td>3.5 ± 0.6</td>
</tr>
<tr>
<td>2-6</td>
<td>3.4 ± 0.3</td>
</tr>
<tr>
<td>2-8</td>
<td>3.4 ± 0.6</td>
</tr>
<tr>
<td>2-12</td>
<td>3.3 ± 0.5</td>
</tr>
</tbody>
</table>

$^b$The data presented are the average of three independent experiments. (n = 25 seedlings/experiment for roots and n = 50 seedlings/experiment for hypocotyls.)
seedlings showed greater bending at 10 mM Ca$^{2+}$ than on the no added Ca$^{2+}$ medium, whereas the transgenic hypocotyls showed more bending on the no added Ca$^{2+}$ medium rather than 10 mM Ca$^{2+}$ (data not shown).

Basipetal Auxin Transport in Roots

Basipetal transport of auxin from the root tip has been shown to be important for the gravitropic response of Arabidopsis roots (Rashotte et al., 2000). Therefore, we compared basipetal auxin transport in vertically grown root tips of wild-type and transgenic seedlings. In three independent experiments, there was a small, but significant, reduction in basipetal transport in the transgenic line 2-8 compared with the wild type (Table V). In both wild-type and transgenic lines, basipetal transport was similarly inhibited by naphthylphthalamic acid (NPA), suggesting that there are no changes in regulation of auxin transport through its NPA-dependent regulatory mechanisms. Rather, these differences are consistent with a reduction in IAA transport capacity. These results suggest that InsP$_3$-mediated signaling affects basipetal auxin transport. The reduction in basipetal transport could contribute to the reduced gravitropic response in the transgenic roots.

Asymmetric Auxin-Induced GUS Expression as a Result of Gravistimulation

The redistribution of auxin has been shown to precede differential growth and the gravity response of plant shoots and roots (Parker and Briggs, 1990; Young et al., 1990). More recently, the lateral redistribution of auxin in gravistimulated roots has been visualized indirectly using the auxin-responsive reporter construct DR5-GUS, as described previously (Rashotte et al., 2001; Buer and Muday, 2004; Hou et al., 2004). We therefore generated transgenic InsP$_5$-ptase lines expressing DR5-GUS (DR5-GUS InsP$_5$-ptase). We first compared GUS staining in vertically grown roots. Both control DR5-GUS and DR5-GUS InsP$_5$-ptase roots showed similar intense staining in the root cap region. We next compared the development of asymmetric auxin-induced GUS expression in response to gravistimulation in the DR5-GUS InsP$_5$-ptase roots and the control DR5-GUS line. Four-day-old light-grown seedlings were gravistimulated in the dark for 4 and 6 h and stained overnight for visualization of GUS activity. GUS activity (as evidenced by strong blue staining) was visible in the root tips of gravistimulated roots.

Table IV. Transgenic roots show a slower rate of curvature after initiation of bending

The kinetics of root bending was measured using the Multi-ADAPT software.

<table>
<thead>
<tr>
<th>Plant Line</th>
<th>Initiation of Curvature</th>
<th>Rate of Curvature after Initiation</th>
<th>Vertical Elongation Rate$^b$</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>min</td>
<td>degree min$^{-1}$</td>
<td>$\mu$m min$^{-1}$</td>
</tr>
<tr>
<td>Wild type</td>
<td>124.6 ± 9.2</td>
<td>0.165 ± 0.014</td>
<td>4.3 ± 0.7</td>
</tr>
<tr>
<td>2-8</td>
<td>155.4 ± 17.1</td>
<td>0.087 ± 0.014</td>
<td>5.3 ± 1.7</td>
</tr>
<tr>
<td>$P$ value</td>
<td>0.1168</td>
<td>0.0031</td>
<td>0.5609</td>
</tr>
</tbody>
</table>

$^a$The value is the average of six individual roots ± se. $^b$The root elongation rates were measured in 50-μm segments from the root tip. $^c$Wild-type and 2-8 values were compared by a two-tailed Student's $t$ test and the $P$ values are reported.
seedlings in both the control DR5-GUS and DR5-GUS InsP 5-птase lines. In addition, several roots of both DR5-GUS and DR5-GUS InsP 5-птase lines also showed a streak of blue staining extending from the tip on the lower side of the root (see Supplemental Fig. 1). This asymmetric expression of DR5-GUS was highly correlated with the angle of bending of the roots. In all roots where a gradient was observed, the angle of bending was between 40° and 45°.

Most significantly, the frequency of the differential GUS staining observed at the 4-h time point was higher in the control DR5-GUS line compared with the DR5-GUS InsP 5-птase line (Table VI), and the average angle of bending in the control was also higher than the InsP 5-птase roots (Table VI). By 6 h of gravistimulation, DR5-GUS InsP 5-птase lines began to catch up. The average angle of bending of the transgenic DR5-GUS InsP 5-птase lines at 6 h was approximately 44° and more roots showed the asymmetric GUS staining. (Note that for these experiments the seedlings were not pretreated for 24 h with dark before gravistimulation and therefore, on average, exhibit a faster bending response than the wild-type and transgenic root responses shown in Fig. 3.) The data show that the frequency of the asymmetric GUS expression correlates well with the angle of bending and that the development of the asymmetric auxin-induced GUS expression is delayed in the InsP 5-птase transgenic roots consistent with their delayed bending response.

Bending Response of Inflorescence Stems

Inflorescence stems of Arabidopsis plants exhibit a fast and robust gravitropic response. We compared the gravitropic response of primary inflorescence stems of 6-week-old wild-type and InsP 5-птase transgenic plants, which were usually 6 to 9 cm tall and did not contain any developed siliques. After 45 min of gravistimulation, inflorescence stems of wild-type and vector control plants showed a greater degree of bending compared to transgenic plants (approximately 75° curvature in wild type and vector control and approximately 50° curvature in transgenic plants; Fig. 5A). The kinetics of inflorescence bending were monitored over a 2-h period (Fig. 5B). As can be seen in the graph, no significant bending takes place until approximately 15 to 20 min. Between 20 and 60 min, the wild-type stems reach and overshoot the vertical (90°) mark. Wild-type bending peaks around 90 min and starts to return to vertical. The transgenic stems were slower to bend and responded to a lesser degree than the wild type. By about 120 min, the bending is comparable in all lines.

Figure 4. The gravitropic response of roots of wild-type and transgenic plants are different on high and low Ca2+ concentrations. A, Vertical growth of 4-d-old wild type (Wt), vector control (C2), and two independent InsP 5-птase transgenic lines (2-6 and 2-8) was monitored after 24 h in the dark on either medium containing no added Ca2+ or medium containing 10 mM Ca2+. B and C, Gravitropic bending of 4-d-old wild type (Wt), vector control (C2), and two independent transgenic lines (2-6 and 2-8) was monitored after 6 (B) and 24 h (C) of gravistimulation in the dark on either medium containing no added Ca2+ (white bars) or medium containing 10 mM Ca2+ (gray bars). Bending is plotted as the average from two independent experiments ± SD (n = 40 roots/experiment). There were statistically significant differences in the bending of each line between the no added Ca2+ versus the 10 mM Ca2+ medium (P < 0.1 using a Student’s t test).
InsP$_3$-Mediated Signaling and Plant Gravitropism

**Table V. Basipetal IAA transport is reduced in the transgenic roots**

<table>
<thead>
<tr>
<th>Plant Line</th>
<th>Basipetal [³H]IAA Transport$^a$</th>
<th>Percentage of Inhibition by NPA</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>−NPA</td>
<td>+NPA</td>
</tr>
<tr>
<td>Wild type</td>
<td>4.2 ± 0.4</td>
<td>2.4 ± 0.2</td>
</tr>
<tr>
<td>2-8</td>
<td>3.1 ± 0.2</td>
<td>1.6 ± 0.1</td>
</tr>
<tr>
<td><em>P</em> value$^b$</td>
<td>0.010</td>
<td>0.00045</td>
</tr>
</tbody>
</table>

$^a$The data presented are the average of two independent experiments ±SD (n = 10 roots/treatment/experiment). $^b$Wild-type and 2-8 values were compared by a two-tailed Student's t test and the _P_ values are reported.

**Table VI. The development of the asymmetric auxin-induced GUS expression is delayed in the InsP 5-ptase transgenic roots compared with the DR5-GUS control**

Histochemical assays and root bending were carried out on DR5-GUS and DR5-GUS InsP 5-ptase roots after 4 and 6 h of bending.

<table>
<thead>
<tr>
<th>Time of Bending Lines</th>
<th>4 h</th>
<th>6 h</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>DR5-GUS</td>
<td>DR5-GUS InsP 5-ptase</td>
</tr>
<tr>
<td>Percent of roots showing gradient$^a$</td>
<td>66%</td>
<td>39%</td>
</tr>
<tr>
<td>Average angle of bending$^b$</td>
<td>41.9 ± 3.8</td>
<td>34.7 ± 2.1</td>
</tr>
</tbody>
</table>

$^a$The data presented are the average of two independent experiments (n = 100 roots/line/experiment). $^b$The so was 1% and 6% for the DR5-GUS and DR5-GUS InsP 5-ptase lines, respectively. $^c$The data presented are the average of three independent experiments ±SD (n = 12 roots/line/experiment).

**DISCUSSION**

The PI pathway is implicated in plant responses to various biotic and abiotic stresses (Stevenson et al., 2000; Meijer and Munnik, 2003). Rapid changes in InsP$_3$ have been documented in different plant systems.
with many different stimuli, including osmotic and cold shock, salinity, light, fungal elicitors, ABA, and gravity. However, it is still not well understood what downstream molecular targets are directly affected by the changes in InsP$_3$ or how these changes lead to a physiological response. To delineate the role of InsP$_3$ in plant responses to stress, we have generated transgenic Arabidopsis plants constitutively expressing the human type I InsP$_{5}$-ptase, an enzyme that specifically hydrolyzes InsP$_3$ (Laxminarayan et al., 1993). The transgenic plants have attenuated InsP$_3$ signaling and therefore are a model system to investigate InsP$_3$-mediated responses. In this article, we test the hypothesis that InsP$_3$ is an integral component of plant gravisignaling.

We first monitored expression of the transgene and basal InsP$_3$ levels in the transgenic plants. The InsP$_{5}$-ptase transgene is stably expressed and the protein is detectable in all tissues tested. Furthermore, in all transgenic tissues tested, basal InsP$_3$ levels were reduced by >90% compared with wild-type levels (Table II). Despite the drastic reduction in basal InsP$_3$ levels, transgenic InsP$_5$-ptase plants showed no obvious morphological differences compared to wild-type plants under normal growth conditions. We have monitored the growth and life cycle of at least three independent lines and all parameters investigated were virtually indistinguishable from wild type (Figs. 1 and 2; Table I).

Of particular interest for the gravitropism studies, the InsP$_5$-ptase transgenic lines showed no defects in elongation growth of the roots or hypocotyls (Table III). However, significantly, the transgenic plants showed a reduced reorientation response upon gravistimulation. This is in contrast to many of the auxin response mutants with altered gravitropism, such as axr3 (Leyser et al., 1996) and axr4 (Hobbie and Estelle, 1995), which have an overall reduced growth phenotype compared to wild type. The data imply that the reorientation response is affected specifically in the InsP$_5$-ptase transgenic plants and that the increase in InsP$_3$ is one component of gravisignaling. When the root-bending response was monitored more closely using Multi-ADAPT software, it revealed that the transgenic roots have a significantly slower rate of curvature after bending is initiated, leading to a reduced response. In addition, our results suggest that InsP$_3$ levels may modulate polar and lateral IAA transport. Measurement of basipetal [$^3$H]IAA movement in vertical roots indicates that transport in the InsP$_5$-ptase transgenic lines is reduced relative to the wild-type plants (Table V). Similar reductions in basipetal transport were previously reported for plants with the eir1/pin2/agr1/wav6 and aux1 mutations (Rashotte et al., 2000, 2001), which alter root basipetal IAA transport and abolish root gravitropism.

![Figure 5](image-url)

**Figure 5.** Inflorescence stems of transgenic plants show a reduced gravitropic bending. A, Gravitropic bending was measured in inflorescence stems of 6-week-old Arabidopsis plants. Plants were first placed upright in the dark for 1 to 2 h and then placed horizontally in the dark. Bending was measured after 45 min. Data plotted are the average of five independent experiments ±SE (n = 25/experiment). B, Gravitropic bending of the inflorescence stems was monitored every 15 min for 2 h. The graph shows the average of six representative wild-type and transgenic inflorescence stems ±SE. The experiment was repeated twice with similar results.

![Figure 6](image-url)

**Figure 6.** Biphasic changes in InsP$_3$ with gravistimulation of inflorescence stems. InsP$_3$ levels were measured in inflorescence stems over a time course of gravistimulation. Data plotted are the average of four independent experiments ±SE (n = 20–30 stems/time point/experiment).
Gravitropism. It should be stressed, however, that there are significant differences between the InsP 5-ptase roots and the eir1/pin2/agr1/wav6 mutants. These mutants show a severe agravitropic phenotype along with altered root elongation and an absence of PIN2 protein in the cortical and epidermal cell files of the root (Luschnig et al., 1998; Muller et al., 1998). In contrast, vertical InsP 5-ptase roots have normal root elongation (Table III), and immunolocalization studies reveal normal PIN2 protein distribution compared to wild-type roots (I.A. Paponov and K. Palme, unpublished data). Furthermore, the gravitropic response is delayed, but not absent, in the InsP 5-ptase transgenic, and, usually, after 48 h root bending approaches wild-type plants.

The asymmetric IAA-induced GUS expression at the root tip after gravistimulation was also delayed in the transgenic plants (Table VI). This delayed formation of the GUS asymmetry on the lower side of the gravistimulated roots is consistent with the reduced levels of basipetal IAA transport, making it more difficult for a gradient of IAA to form. The timing of formation of the gradient is still linked to the curvature of the roots, suggesting that the delay in gravity response is linked to the delayed formation of a lateral gradient of IAA. These results suggest that InsP3 may be involved in establishing the timing of the response. Taken together, the fact that basipetal IAA transport was reduced along with the delay in lateral redistribution of auxin-induced GUS expression in the InsP 5-ptase roots with gravistimulation, these data imply that the InsP3-mediated signaling events must occur upstream of polar and lateral auxin transport and that attenuation of InsP3-mediated signaling either directly or indirectly affects the dynamics of auxin translocation.

Although many plant tissues respond to gravity, there are significant differences in how and where these responses are mediated. There are some basic differences in the architecture of roots versus shoots and also in the site of gravity sensing and gravity response in different organs. For example, in plant roots, the columella cells of the root cap are involved in gravity sensing, while the differential growth response takes place primarily in the distal elongation zone of the root (Ishikawa and Evans, 1993; Rosen et al., 1999). In cereal grasses such as oats and maize, both gravity sensing and response take place in a specialized tissue in the stem known as the pulvinus (Kaufman et al., 1995). In herbaceous dicot hypocotyls and floral stems, the gravitropic response occurs throughout the entire elongation zone of the stem (Fukaki et al., 1996a, 1996b). The mechanisms of gravitropism in roots and shoots therefore may have both common and separate steps. The differences are illustrated by some of the altered gravity mutants, which show a gravitropic defect in only a subset of the gravityresponding organs. For example, the sgr mutants, which are defective in hypocotyls and/or inflorescence stem gravitropism, show a normal gravitropic response in the roots (Yamauchi et al., 1997; Kato et al., 2002; Yano et al., 2003). Alternatively, several of the auxin response (aux1, axr3, and axr4) mutants have normal hypocotyl and inflorescence bending, although root gravitropism is impaired (Hobbie and Estelle, 1995; Leyser et al., 1996; Marchant et al., 1999). In contrast, the InsP 5-ptase transgenic plants show a reduced gravitropic response in roots, hypocotyls, and inflorescence stems.
The transgenic response is slower and reduced by 30% compared with the wild type. These results support the idea that InsP$_3$ is a fundamental conserved component in plant gravisignaling. However, it is clear that multiple parallel or converging signaling pathways must be involved, since the gravitropic response is not completely eliminated in the transgenic plants.

In previous work, we demonstrated that biphasic increases in InsP$_3$ are associated with the gravitropic responses of maize and oat pulvini (Perera et al., 1999, 2001); however, until now, it was not clear whether InsP$_3$ signaling was a unique feature of the pulvinus system. Using inflorescence stems of Arabidopsis plants, we have quantified the levels of InsP$_3$ during gravisimulation. Although it was not feasible to separate the inflorescence stems into upper and lower sides, we have detected biphasic increases in InsP$_3$ in the total stems with gravisimulation. These InsP$_3$ changes are similar to the pattern seen with the lower sides of the pulvini and indicate that InsP$_3$ increases with gravisimulation in both monocots and dicots. Taken together with the fact that the transgenic plants with attenuated InsP$_3$ signaling exhibit reduced gravitropic responses, these results are strong evidence for a universal role for polyphosphoinositides and the generation of InsP$_3$ in the gravity signal transduction cascade in plants.

The involvement of InsP$_3$ in gravitropism also implicates calcium. There is much indirect evidence supporting a role for calcium in gravisignaling and response (for review, see Sinclair and Trewavas, 1997; Fasano et al., 2002). However, it has been difficult to directly measure Ca$^{2+}$ changes in plants in response to gravisimulation. Legue et al. (1997) carried out a careful study of Arabidopsis root tips and were unable to detect changes in Ca$^{2+}$ upon gravisimulation. More recently, Plieth and Trewavas (2002) have shown a global Ca$^{2+}$ signal upon reorientation by monitoring the luminescence of a large population of Arabidopsis seedlings expressing aequorin. The signature of the Ca$^{2+}$ transient with gravisimulation consisted of a rapid spike followed by a longer shoulder lasting about 15 min. Interestingly, mechanical stimulation resulted in only a quick spike as did wind and touch. Based on the results that show that the shoulder exhibits signal adaptation (i.e. the amplitude of the shoulder was affected by the degree of reorientation and also by the number of successive reorientations), whereas the spike was unchanged, the authors infer that the spike and shoulder might have different cellular or subcellular origins (Plieth and Trewavas, 2002).

The biphasic patterns of Ca$^{2+}$ signals revealed with aequorin are reminiscent of the pattern of InsP$_3$ changes with gravisimulation (Perera et al., 1999, 2001). The initial transient fluctuations in InsP$_3$ detected within seconds of gravisimulation were proposed to be part of a wake-up call, which alerts the plant to a change in its environment (Perera et al., 2001). These initial transients are consistent with the rapid, transient aequorin-sensitive Ca$^{2+}$ signal and may be common to many different stimuli. The longer sustained increase in InsP$_3$, which only occurs on the lower side of the pulvinus during gravisimulation, precedes and correlates with the bending response and may be more specific for the gravistimulus. This would be analogous to the longer Ca$^{2+}$ shoulder.

In the classical signal transduction scheme, a localized change in Ca$^{2+}$ is propagated through the cell and to neighboring cells via the soluble second messenger InsP$_3$ (Tucker and Boss, 1996). In the transgenic cells, InsP$_3$ is constantly being degraded by the InsP$_3$ 5-tpase enzyme. Therefore, any InsP$_3$-mediated Ca$^{2+}$ signal generated in response to a stimulus would not spread throughout the cell and to neighboring cells. Additionally, since InsP$_3$-mediated Ca$^{2+}$ signaling is attenuated, the InsP$_3$-sensitive intracellular Ca$^{2+}$ stores may always be available in the transgenic plants. This may help explain why the transgenic lines showed a stronger reorientation response on the no added Ca$^{2+}$ compared to the 10 mM Ca$^{2+}$ medium.

The cold gravisimulation experiments have also been illuminating. These experiments are based on the fact that plants can sense a gravistimulus in the cold and respond to that stimulus when returned to room temperature. Fukaki et al. (1996a) showed that the memory or stored signal of the cold gravistimulus is retained for up to 60 min in Arabidopsis inflorescence stems. Wyatt et al. (2002) have utilized this phenomenon to devise a creative screen for mutants impaired in some aspect of gravity signaling. We have shown previously that increases in InsP$_3$ can occur at 4°C in gravisimulated oat pulvini (Perera et al., 2001). Furthermore, since the sedimentation of statoliths is not impaired by cold temperature (Wyatt et al., 2002), it is clear that gravity sensing and some of the early signaling can occur in the cold. In contrast, auxin transport is dramatically decreased in the cold (Morris, 1979; Wyatt et al., 2002), and cold temperature also affects cytoskeleton and membrane dynamics (Mizuno, 1992; Örvar et al., 2000; Sangwan et al., 2001).

When wild-type Arabidopsis plants were gravisimulated in the cold and returned to room temperature, within the first 15 to 30 min there was a rapid positive response to the cold gravistimulus, which was followed by a return to the vertical. In some instances, the wild-type inflorescence stems display an overshooting response (bending beyond the vertical) following the return to room temperature. In contrast, the transgenic plants show a dampened response, and the rate of the bending response is significantly slower compared to wild type. In fact, the transgenic response to gravisimulation in the cold is reduced by 50% to 60% compared with the wild type, which is a greater reduction than seen when plants were gravisimulated at room temperature. These data imply that the contribution of InsP$_3$-mediated signaling is magnified in the cold, possibly because other signaling mechanisms are inhibited by low temperature.

An important feature of the InsP 5-tpase transgenic plants is that, although InsP$_3$ does not accumulate to wild-type levels, InsP$_3$ synthesis via PLC is not
inhibited (Perera et al., 2002), and, therefore, there is no buildup of lipid precursor (PtdInsP₂) or a block in phosphatic acid production via diacylglycerol (DAG). The expression of the highly active recombiant InsP₅-5-ptase results in the rapid hydrolysis of InsP₅ and possibly an increased flux through the PI pathway. However, the transgenic plants grow normally and show no obvious phenotype under normal conditions. In contrast, a PI PLC knockout (plc1) in the moss Physcomitrella, while exhibiting an altered gravitropic response, also shows several pleiotropic growth defects, such as reduced gametophore formation, reduced levels of chlorophyll, and insensitivity to cytokinin (Repp et al., 2004). This is not surprising, as a loss of PLC activity would lead to reduced InsP₅ levels as well as reduced DAG and DAG-mediated phosphatic acid levels and would also affect PtdInsP₂ accumulation. We suspect that, for these reasons, using antisense or knockout strategies of the PLC enzymes will have profound effects on plant growth beyond affecting InsP₅ levels. Because the InsP₅-5-ptase transgenic Arabidopsis plants have altered InsP₅ turnover and a dampened InsP₃ signal, but no blockage in the pathway, they provide a good model system for specifically analyzing downstream InsP₅-mediated responses.

In summary, we have shown that dampening InsP₅-mediated signaling delays the timing and reduces the magnitude of the gravitropic response of Arabidopsis roots, hypocotyls, and inflorescence stems, and we propose that InsP₅ is a fundamental component of plant gravisignaling that is upstream of auxin redistribution. Future work will focus on understanding the interaction of the InsP₃ signalling pathway with other early components of the gravity signal transduction cascade.

MATERIALS AND METHODS

Plant Transformation and Selection of Transgenic Lines

The cDNA encoding the human type I InsP₅-5-ptase (accession no. X77567) was subcloned into the Xhol site of the pKYL71-35S<sup>Ⅱ</sup> binary vector (Schardl et al., 1987). The pKYLI-35S<sup>Ⅱ</sup> vector contains a modified 35S promoter, 3′-untranslated region of the pea (Pisum sativum) small subunit of Rubisco E-9 gene and a plant kanamycin resistance cassette. The coding region of the InsP₅-5-ptase gene, along with the 5′-end His tag, was amplified from the bacterial expression vector pQE31 using forward and reverse primers engineered to contain Xhol sites. The orientation of the resulting plasmid, pKYLI-35S<sup>Ⅱ</sup>-InsP₅-5-ptase, was verified by restriction enzyme analysis and DNA sequencing. The binary plasmids pKYLI-35S<sup>Ⅱ</sup>-InsP₅-5-ptase and pKYLI-35S<sup>Ⅱ</sup>-vector control were electropropated into Agrobacterium tumefaciens strain GV<sub>2000</sub> (Ishikawa and Evans, 1997) as described by Rashotte et al. (2001). A mixture containing 1% agar, 5mM MES buffer/1% (w/v) Suc, MS salts, 1% Suc, MES buffer, pH 5.7, and 0.8% type M agar. Plates were incubated vertically in a growth chamber under short-day conditions (8 h light/16 h dark) at 21°C with light intensity of approximately 150 μmol m<sup>⁻²</sup> s<sup>⁻¹</sup>. For root and hypocytol elongation measurements, 4 d after germination plates were covered and placed in the dark and growth was monitored every 24 h for a 3- to 4-d period. For morphometric analysis, surface-sterilized seeds were sown on soil (PGX soil mix; Hummert), incubated in a growth chamber under long-day conditions (16 h light/8 h dark), and four to six plants of each line were monitored over an 8-week period as described by Boyes et al. (2001). Briefly, every 2 to 3 d plants were observed and leaf number, leaf length, and rosette size were measured. Across development, the timing of leaf expansion, emergence of the primary inflorescence stem, flower development, and silique formation were also measured, and seeds were collected to determine yield.

Gravitropic Bending Measurements

For root bending, seeds were germinated as described above and grown on vertically oriented plates for 3 d in the light and 1 d in the dark. Plates were then rotated by 90° and incubated in the dark. Plates were photographed prior to turning and at 2, 6, and 24 h after turning. Images were captured using a Hamamatsu color camera attached to a Leica stereo dissecting microscope or a Nikon Coolpix 4500 digital camera. Photographs were analyzed and bending angles were measured using Adobe Photoshop and analyzed using Microsoft Excel. Root bending was also monitored using Multi-ADAPT software (Ishikawa and Evans, 1997) as described by Rashotte et al. (2001).

For hypocotyl bending, seeds were grown on vertically oriented plates for days in the light followed by 1 d in the dark. Plates were then rotated by 90° and incubated in the dark for 48 h. Images were captured prior to turning the plates and 24 and 48 h after turning using a Hamamatsu color camera attached to a Leica stereo dissecting microscope. Bending angles were measured using Adobe Photoshop and analyzed using Microsoft Excel.

For inflorescence bending, Arabidopsis plants were grown in pots (20 seeds/pot) for 4 weeks under short-day conditions followed by 2 weeks under long-day conditions. Experiments were carried out when primary inflorescence stems were 6 to 9 cm long. Plants were first incubated in the dark for 2 h prior to turning and then the pots were oriented horizontally in the dark. Bending was monitored every 15 min for 2 h using a Nikon Coolpix 4500 digital camera. For cold bending experiments, plants were incubated horizontally in the dark at 4°C for 1 h. Plants were then returned to room temperature and placed vertically in the dark and monitored every 15 min for 2 h.

Statistical Analysis

For all growth and bending measurements, data were subjected to statistical analysis by one-way ANOVA. The mean values for each line were compared using Student’s t test assuming equal variance.

Root Basipetal Transport Assays

Root auxin transport measurements were made on 5-d-old vertically grown seedlings, using a previously published procedure (Rashotte et al., 2000, 2001). A mixture containing 1% agar, 5 mM MES buffer/1% (w/v) Suc, pH 5.5, and 100 nM 3H-HA was prepared in a 3-mL scintillation vial. A narrow stem transfer pipette was carefully inserted into the hardened agar such that a

(Weigel and Glazebrook, 2002). DNA was digested by restriction enzymes and analyzed by gel electrophoresis, followed by Southern blotting, using the InsP₅-5-ptase gene as a probe. Four independent transformed lines were further selected for two more generations. Stable expression of the transgene was monitored by immunoblotting as described below.
Histochemical Assays for Expression of the Auxin Reporter DR5-GUS

InsP$_3$ Assays

Root and shoot samples were harvested from 5-d-old seedlings grown on plates and frozen immediately in liquid N$_2$. For inflorescence-bending time courses, inflorescence stems (15–20 time point) were harvested without any flowers and siliques and frozen immediately in liquid N$_2$. Frozen tissue (approximately 0.05 g) was ground to powder in liquid N$_2$ and incubated with 150 µl of 10% perchloric acid on ice for 15 min. Samples were centrifuged to remove the precipitate and the supernatant was transferred to a new tube and the pH adjusted to 7.5 using 1.5 M KOH/3 M HEPES. InsP$_3$ assays were carried out using the TRK1000 InsP$_3$ assay kit (Amerham-Pharmacia Biotech) as described previously (Perera et al., 1999, 2001, 2002).

Protein Isolation and Immunoblotting

Root and shoot samples were harvested from 5-d-old Arabidopsis seedlings grown on plates and frozen immediately in liquid N$_2$. For inflorescence-bending time courses, inflorescence stems (15–20 time point) were harvested without any flowers and siliques and frozen immediately in liquid N$_2$. Frozen tissue (approximately 0.05 g) was ground to powder in liquid N$_2$ and incubated with 150 µl of 10% perchloric acid on ice for 15 min. Samples were centrifuged to remove the precipitate and the supernatant was transferred to a new tube and the pH adjusted to 7.5 using 1.5 M KOH/3 M HEPES. InsP$_3$ assays were carried out using the TRK1000 InsP$_3$ assay kit (Amerham-Pharmacia Biotech) as described previously (Perera et al., 1999, 2001, 2002).

Sequence data from this article can be found in the GenBank/EMBL data libraries under accession number X77567.

ACKNOWLEDGMENTS

We thank Dr. Sarah E. Wyatt (Ohio University, Athens, OH) and Dr. John Love (University of Exeter, UK) for advice with the plant transformation and selection of primary transformants. We acknowledge Matthew Keefe, Beth Stapperferne, Katie Kovac, and Candace Randall for their hard work and help in selecting and characterizing the transformants and carrying out the growth studies, and Kelly Altbaus for carrying out the GUS histochemical assays. Thanks to “Jeff” Yue Xu, Dr. Eva Johannes, and Dr. Nina Allen of the Cell and Molecular Imaging Facility at North Carolina State University for help with microscopy, Dr. Linda Hanley-Bowdoin (North Carolina State University) for use of the dissecting microscope, and Peter Aspes Jr. for the digital photography. We also thank Dr. Ingo Heilmann (University of Göttingen, Germany) and the members of the Boss lab (North Carolina State University) for helpful suggestions and discussion.

Received December 5, 2005; revised December 5, 2005; accepted December 13, 2005; published December 29, 2005.

LITERATURE CITED


Received December 5, 2005; revised December 5, 2005; accepted December 13, 2005; published December 29, 2005.


Copyright © 2006 American Society of Plant Biologists. All rights reserved.


