Transformation of shoots into roots in Arabidopsis embryos mutant at the TOPELESS locus

Jeff A. Long1,3, Scott Woody1, Scott Poethig2, Elliot M. Meyerowitz3 and M. Kathryn Barton1,*,†

1Department of Genetics and Program in Cellular and Molecular Biology, University of Wisconsin-Madison, Madison, WI, USA
2Department of Biology, University of Pennsylvania, Philadelphia, PA, USA
3Division of Biology, California Institute of Technology, Pasadena, CA, USA

*Present address: Department of Plant Biology, Carnegie Institution of Washington, 260 Panama Street, Stanford, CA 94305, USA
†Author for correspondence (e-mail: barton@andrew2.stanford.edu)

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SUMMARY

We describe a novel phenotype in Arabidopsis embryos homozygous for the temperature-sensitive topless-1 mutation. This mutation causes the transformation of the shoot pole into a root. Developing topless embryos fail to express markers for the shoot apical meristem (SHOOT MERISTEMLESS and UNUSUAL FLORAL ORGANS) and the hypocotyl (KNAT1). By contrast, the pattern of expression of root markers is either duplicated (LENNY, J1092) or expanded (SCARECROW). Shifts of developing topless embryos between permissive and restrictive temperatures show that apical fates (cotyledons plus shoot apical meristem) can be transformed to basal fates (root) as late as transition stage. As the apical pole of transition stage embryos shows both morphological and molecular characteristics of shoot development, this demonstrates that the topless 1 mutation is capable of causing structures specified as shoot to be respecified as root. Finally, our experiments fail to show a clear link between auxin signal transduction and topless-1 mutant activity; the development of the apical root in topless mutant individuals is not dependent on the activity of the predicted auxin response factor MONOPTEROS nor is the expression of DR5, a proposed ‘auxin maximum reporter’, expanded in the apical domain of topless embryos.

Key words: Arabidopsis, TOPELESS, Shoot transformation, Embryogenesis, Polarity

INTRODUCTION

Shoot and root systems of land plants carry out complementary functions fundamental to plant survival. These systems form from meristems established at opposite poles of the embryo. Morphological indications of differences along the shoot-root axis are most clearly manifested after the globular stage of embryogenesis. Globular stage embryos consist of two parts: a spherical mass of cells (the embryo proper) that gives rise to the seedling plant, and a suspensor, often filamentous, that anchors the embryo proper to the ovule and may act as a conduit for nutrients from the mother plant. In Arabidopsis, the suspensor consists of a single row of cells giving the globular stage embryo the appearance of a lollipop. In Arabidopsis, as in many other dicotyledonous plant species, the future shoot-root axis of the globular embryo proper is aligned parallel to the long axis of the suspensor. The root pole forms from the side of the globular embryo closest to the suspensor, while the shoot pole forms from the side directly opposite. As the embryo makes the transition from the globular to the heart stage of embryogenesis, growth occurs more rapidly in the apical-basal dimension than in the radial dimension. Concomitant with this, morphological features of the shoot (cotyledons and shoot apical meristem) and root poles (organization of cell files and root cap) develop.

Polarity along the long axis of the globular stage embryo is foreshadowed in the polar organization of the egg cell and the zygote (Mansfield and Briarty, 1991; Mansfield et al., 1991). The egg cell is positioned in the embryo sac with the basal end facing the micropyle (the opening through which the pollen tube enters) and the apical end facing the central cell. The egg cell itself is polarized with the nucleus located closer to the apical end and the vacuole located closer to the basal end. The first division of the zygote is unequal, generating a large basal cell and a smaller terminal cell. The larger basal cell gives rise to the suspensor, while the smaller apical cell generates the embryo proper. The basal cell also gives rise to the hypophysis. The hypophysis is the only descendent of the basal cell to generate a portion of the seedling; it gives rise to portions of the root meristem and root cap (Mansfield and Briarty, 1991; Dolan et al., 1993).

Mutations in the GNOM/EMB30 gene disrupt both the polarity of the zygote and of the developing embryo proper (Mayer et al., 1993). The orientation of the first zygotic cell division is randomized in gn mutant embryos and the most severely affected embryos develop as green balls that lack an axis. Polarity appears to be abolished at the cellular level in gn
mutants. Consistent with this, the PIN1 molecule, which is normally localized to the basal pole of the cell, is found around the circumference of the cell or localized to a side of the cell that bears no relationship to the apical-basal axis (Steinmann et al., 1999). The GN/EMB30 gene encodes a GEF/ARF-like molecule that may mediate its effects through targeted secretion (Shevell et al., 1994; Busch et al., 1996).

While polar cues embedded in the egg cell or early embryo may be important for early development of the embryo and for the proper alignment of the embryonic axis with that of the embryo sac, such cues are not absolutely required for the development of an embryo with a shoot-root axis. Ectopic embryos form from cells of the suspensor in some mutants (Vernon and Meinke, 1994). Embryos that form in this way may be oriented in the same or in the opposite direction to the embryo from which they derive. In addition, somatic embryos with a normal shoot-root axis can develop from a variety of tissue types.

Several mutants exist in which the development of the shoot or the root pole is disrupted, but in which a polar axis is nonetheless generated. For example, monopterous, bodenlos and axr6 mutations all interfere with the development of the root pole (Hardtke and Berleth, 1998; Hamann et al., 1999; Hobbie et al., 2000). Conversely, cup-shaped cotyledon (cuc) and shoot meristemless (stm) mutations interfere with the development of the cotyledons and meristem (Barton and Poethig, 1993; Aida et al., 1997). Such genes presumably act downstream of a signal that distinguishes cells of the future root pole from cells of the future shoot pole.

In cultured cells, ratios of the plant growth regulators cytokinin and auxin have been shown to play a role in the process of specifying cells as root or shoot (Skoog and Miller, 1957). High auxin to cytokinin ratios promote root development, while high cytokinin to auxin ratios promote shoot development. It is still unknown whether such ratios play a role in the embryo. However, the observation that axr6 mutants fail to develop a root and also fail to respond normally to auxin suggests they may (Hobbie et al., 2000).

One mutation, doppelwurzel, may cause a polar duplication of pattern during embryogenesis as it causes two complete seedlings to develop from a single ovule in a head to head (cotyledon to cotyledon) configuration (Jürgens et al., 1991). However, little is known about embryonic development in this mutant and it is possible that the apparent polar duplication is actually the result of the development of twin embryos from a single zygote.

We describe here a novel phenotype caused by the topless-1 (tpl-1) mutation. In homozygous tpl-1 mutants the apical pole of the embryo develops as a root instead of as a shoot. The result is a seedling with two roots joined at the base and no hypocotyl, cotyledons or shoot apical meristem. The novel tpl-1 mutation should provide an entrance to molecular genetic study of this important, but previously experimentally intractable, decision in plant development.

**MATERIALS AND METHODS**

**Genetics**

The tpl-1 mutation was isolated in the Landsberg erecta genetic background after EMS mutagenesis. Three backcrosses to Landsberg erecta were performed prior to phenotypic analysis. The tpl-1 mutation was mapped using CAPS markers as described by Konieczny and Ausubel (Konieczny and Ausubel, 1993). tpl-1 mutant individuals were crossed to wild-type Columbia individuals. The F1 progeny were allowed to self-fertilize at 24°C. DNA was prepared from F2 seedlings exhibiting strong topless mutant phenotypes (monocot or more extreme). Three out of 268 chromosomes were recombinant between tpl-1 and CAPS marker g2395 (recombinant frequency=1.1%). Two out of 268 chromosomes were recombinant between tpl-1 and CAPS marker m59 (recombinant frequency=0.8%). In all cases, chromosomes recombinant between one marker and tpl-1 were non-recombinant for the other marker, indicating that tpl-1 lies between them.

**In situ hybridization experiments**

In situ hybridization and probe preparation were performed as in Long and Barton (Long and Barton, 1998). The KNAT1, SCR, STM, ANT and UFO probes have been described by Lincoln et al. (Lincoln et al., 1995), DiLaurenzio et al. (DiLaurenzio et al., 1996), Long et al. (Long et al., 1996), Long and Barton (Long and Barton, 1998) and Lee et al. (Lee et al., 1997), respectively.

**Histology**

Staining of β-glucuronidase (GUS) activity was performed by incubating embryos dissected from their ovules overnight at 37°C in 2 mM 5-bromo-4-chloro-3-indolyl b-D-glucuronic acid, 100 mM NaPO4, pH 7, 0.1% Triton X-100, 1mM FeCN and 10 mM EDTA.

Fig. 1. Representative phenotypes of seedlings homozygous for the tpl-1 mutation. (A) Wild-type seedling. (B) tpl-1 homozygote that has developed a single cotyledon and lacks a shoot apical meristem. (C) tpl-1 homozygotes that has developed a cup-shaped cotyledon and no shoot apical meristem. (D) tpl-1 homozygote that has developed with no detectable cotyledon tissue or shoot apical meristem. (E) tpl-1 homozygote that has developed both an apical (a) and a basal (b) root. Scale bars: 1 mm.
**Table 1. The *topless-1* allele is temperature sensitive**

<table>
<thead>
<tr>
<th>Temperature</th>
<th>One or two cotyledons, no SAM*</th>
<th>Tube-shaped cotyledon, no SAM</th>
<th>SAM and cotyledons absent</th>
<th>Shoot replaced by a root</th>
</tr>
</thead>
<tbody>
<tr>
<td>24°C</td>
<td>57% (n=254)</td>
<td>36% (n=158)</td>
<td>4% (n=17)</td>
<td>3% (n=12)</td>
</tr>
<tr>
<td>28°C</td>
<td>7% (n=25)</td>
<td>16% (n=61)</td>
<td>13% (n=49)</td>
<td>64% (n=246)</td>
</tr>
</tbody>
</table>

*Absence of SAM refers to SAM formed during embryogenesis, not to postembryonic SAM formation. Nearly all seedlings of this class are able to form a SAM within 2 weeks of germination.

**Temperature shift experiments**

Homozygous *tpl-1* plants were grown at 24°C until they began to flower. For each time point, 10-12 *tpl-1* flowers were emasculated and allowed to mature at 24°C for 2 days. Pollen from *tpl-1tpl-1* plants was then crossed onto the carpels and the plants were immediately placed at either 29°C or 17°C. To determine the developmental stage of the embryos being shifted, embryos from one hand pollination were dissected and examined microscopically at the time of each temperature shift.

**RESULTS**

**topless mutant phenotype – seedling stage**

Plants carrying the *topless-1* (*tpl-1*) mutation were identified in a screen for seedlings defective in shoot apical meristem (SAM) formation (Barton and Poethig, 1993). The *tpl-1* mutation maps to the upper arm of chromosome 1 between markers g2395 and m59 (see Materials and Methods for data).

When homozygous, the *tpl-1* mutation causes either loss of apical structures or transformation of the apical pole of the embryo into a root. The *tpl-1* mutation is temperature sensitive (Table 1). Lower temperatures increase the fraction of seedlings with cotyledon tissue and decrease the fraction of double-root seedlings. Conversely, higher temperatures increase the fraction of severely mutant seedlings, especially those with two roots.

Fig. 1B-D shows homozygous *tpl-1* seedlings that fall into the ‘loss’ category. The mildest phenotype is a seedling with two cotyledons (fused to varying extents) and no SAM (not shown). Homozygous *tpl-1* seedlings may also develop a single cotyledon and no shoot apical meristem (Fig. 1B, Fig. 2B). More severely affected seedlings form a radially symmetric, cup-shaped cotyledon and lack a shoot apical meristem (Fig. 1C, Fig. 2C). Yet others appear to lack cotyledons and SAM completely, and only make a hypocotyl (Fig. 1D, Fig. 2D). (It is because of this last phenotype that the gene was named *topless*.)

Fig. 1E and Fig. 2E-F show homozygous *tpl-1* seedlings that fall into the ‘transformation’ category. In these seedlings, an apical root replaces the shoot. The apical root has a root cap and exhibits normal gravitropism. The cellular organization of the apical root is roughly similar to a wild-type root with long files of cells converging near the root tip (Fig. 3). However, the apical root is often broader and appears to incorporate more cells than the wild-type basal root (Fig. 2F, Fig. 3).

Rarely, homozygous *tpl-1* seedlings show simultaneous expression of apical and basal characteristics. Fig. 2G shows a homozygous *tpl-1* seedling in which root hairs grow from a reduced cotyledon.

A fraction of *topless* mutant individuals are able to form a SAM by a few weeks after germination. These SAMs form at the adaxial base of the cotyledon. Such shoot apical meristems go on to form fairly normal, fertile plants which serve as a source of homozygous *tpl-1* seed. The most penetrant abnormality detected in such *tpl-1* ‘escape’ plants is a tendency for the carpels to be unfused (not shown). Nearly all *tpl* seedlings with one or two cotyledons are able to form SAMs postembryonically. *topless* mutants with cup-shaped cotyledons do so only rarely, while *tpl-1* mutants in which the apical pole is transformed into a root have never been observed to form a SAM.

Seedlings heterozygous for *tpl-1* sometimes exhibit mild fusion of the cotyledons. To determine whether this is due to a dominant, zygotic effect of *tpl-1* or to a maternal effect of...
the tpl-1 mutation, reciprocal crosses were carried out. The fraction of heterozygotes showing a cotyledon fusion defect was the same whether the maternal or paternal parent was homozygous for tpl-1 (Table 2). Thus, there is no indication of a maternal effect in these experiments. Instead, we conclude the weak cotyledon fusion defect in tpl-1 heterozygotes is dependent on the genotype of the zygote and thus due to weak dominance of the tpl-1 allele.

### Development of topless mutant embryos

tpl-1 mutant embryos are initially morphologically identical to wild-type embryos. At the globular stage they are similar to wild-type embryos in size, shape and cellular organization (see Fig. 5J). By the heart stage, tpl-1 mutant embryos differ from wild-type embryos. They are oblong and lack apparent cotyledon primordia (Fig. 4). The oblong tpl embryos continue to elongate during embryogenesis, accumulate storage proteins and mature.

To further determine the extent to which tpl mutants deviate from wild-type embryos in early development, a panel of markers for apical and basal fates was examined in mutant embryos.

### Markers associated with apical fates

Mutants homozygous for tpl-1 fail to form a SAM and frequently also lack cotyledons. These defects could be due to a failure to specify cells as SAM or cotyledon cells in the apical region of the globular embryo. Alternatively, they could be due to a failure of properly specified cells to develop into recognizable structures. To distinguish between these possibilities, we analyzed markers normally expressed in wild-type embryos with fused cotyledons in topless mutant embryos.

The **SHOOTMERISTEMLESS** (STM) gene is required for initiation and maintenance of the SAM (Barton and Poethig, 1993; Long and Barton, 1998). In wild-type embryos, STM RNA marks the developing SAM and is first detected at the late globular stage, before there is morphological evidence of a shoot apical meristem. In homozygous tpl-1 embryos, the STM transcript is not detected (n>50; Fig. 5B). Wild-type siblings segregating in the same silique and processed along with the tpl-1 embryos exhibited normal STM expression (Fig. 5A). tpl-1 embryos accumulate transcripts related to cellular functions, such as the UBQUITIN transcript, which could be detected throughout the tpl-1 embryo (Fig. 5D).

**UFO** is also expressed in the SAM during embryogenesis. Accumulation of the UFO transcript is dependent on wild-type STM function (Long and Barton, 1998). Given the failure of STM expression in tpl-1 mutants, we would expect UFO transcript also to be missing. Indeed, UFO mRNA fails to accumulate in tpl-1 mutants (n>50; Fig. 5C). As in the STM experiment, control wild-type embryos processed simultaneously showed normal UFO expression (not shown).

Finally, the distribution of a transcript ordinarily found in the developing hypocotyl was examined. **KNAT1** is normally expressed in peripheral regions of the hypocotyl (Fig. 5G). In tpl-1 mutants, KNAT1 transcript is either missing or only faintly expressed (Fig. 5H). Loss of KNAT1 transcript accumulation is somewhat surprising as tpl-1 mutant seedlings grown at 24°C usually make a structure recognizable as a hypocotyl. The expression data suggest that the program for hypocotyl development may be somewhat impaired even at this relatively permissive temperature.

The above experiments were done at 24°C, a temperature that does not cause a particularly large number of double root seedlings (Table 1). The experiments have not been carried out

### Table 2. The tpl1 mutation is incompletely dominant

<table>
<thead>
<tr>
<th>Temperature (°C)</th>
<th>Female parent</th>
<th>Male parent</th>
<th>Number of progeny with fused cotyledons</th>
</tr>
</thead>
<tbody>
<tr>
<td>29</td>
<td>TPL/TPL</td>
<td>.tpl/tpl</td>
<td>86/390 (22%)</td>
</tr>
<tr>
<td></td>
<td>tpl/tpl</td>
<td>TPL/TPL</td>
<td>100/429 (23%)</td>
</tr>
<tr>
<td>24</td>
<td>TPL/TPL</td>
<td>.tpl/tpl</td>
<td>4/195 (2.1%)</td>
</tr>
<tr>
<td></td>
<td>tpl/tpl</td>
<td>TPL/TPL</td>
<td>3/178 (1.7%)</td>
</tr>
<tr>
<td>17</td>
<td>TPL/TPL</td>
<td>tpl/tpl</td>
<td>2/144 (1.4%)</td>
</tr>
<tr>
<td></td>
<td>tpl/tpl</td>
<td>TPL/TPL</td>
<td>3/135 (2.2%)</td>
</tr>
</tbody>
</table>
at 29°C. However, given that there is even less development of apical structures at 29°C, it is unlikely that expression of STM, UFO or KNAT1 would be restored at the higher temperature.

The experiments with the UFO, STM and KNAT1 markers indicate that the failure of apical structures to develop in tpl-1 mutants is due to a failure to specify apical (shoot) fates, rather than a failure of properly specified domains to grow and develop into morphologically recognizable structures.

The AINTEGUMENTA (ANT) RNA marks developing leaf, floral organ and cotyledon primordia (Elliott et al., 1996; Long and Barton, 1998). In the wild-type embryo ANT is first expressed in the globular stage embryo prior to any morphological evidence of cotyledon initiation. The two cotyledon primordia appear to be specified initially as a single donut-shaped primordium encircling the apical half of the globular embryo (Fig. 5E). In tpl-1 mutants, by contrast, ANT transcript is found in the center of the apical pole of the embryo in addition to the peripheral ring (n>200; Fig. 5F). ANT transcript was also found in topless embryos developing at the most restrictive temperature (29°C). However, at this temperature, it is often confined to a more apical region: the outermost (protodermal) layer (not shown).

The experiments with the ANT marker indicate that some cells are specified as cotyledon in tpl-1 mutants, even at the most restrictive temperature. However, the pattern of cells specified is altered: the altered pattern shows changes both in the apical/basal dimension of the embryo as well as in the radial (or central/peripheral) dimension.

Markers associated with basal fates

Homozygous tpl-1 mutant embryos developing at 29°C show a high frequency of apparent shoot to root transformation. To further investigate the development of such double root embryos, we analyzed the expression of three markers for root cell fates. The SCARECROW transcript accumulates in the lenticular cell and in the cortical cell files (Fig. 5I) (DiLaurenzio et al., 1996). In the wild-type embryo, SCR expression is limited to the region below the O boundary. (The O boundary divides the globular stage embryo into apical and basal halves. It is created when the four cells of the quadrant stage embryo divide to generate the octant stage embryo.) In tpl-1 mutant embryos, scr extends above the O boundary, indicating that root and/or hypocotyl fates extend further apically in topless mutant embryos (Fig. 5J).

The LENNY reporter was isolated in a screen of enhancer trap lines. [The approach used has been described by Sundaresan et al. (Sundaresan et al., 1995).] The LENNY reporter expresses GUS activity in the root pole of the developing embryo and not in the shoot pole (Fig. 6A,B). LENNY is initially expressed in the lenticular cell (a descendent of the hypophysis) beginning at the heart stage. Expression later spreads to encompass surrounding cells in the root apical meristem.

Twenty-one percent (16/76) of homozygous topless mutant embryos developing at 29°C show expression of the LENNY reporter at both poles of the embryo by 168 hours postfertilization (Fig. 6D-F). (Wild-type embryos grown at 29°C are typically at bent cotyledon stage by 168 hours.) The extent to which embryos express the LENNY reporter varies. Some embryos show only a weak area of LENNY expression at the apical pole (Fig. 6D). More commonly, both poles express LENNY with near equal intensity (Fig. 6E,F).

Note that the apical root primordium does not have the same cellular organization as the normal, basal, root primordium. For example, the files of cells that make up the developing vasculature are not apparent in the tpl apical root (Fig. 6E,F). In addition, the apical root primordium is not tapered as in a normal root but instead is much wider.

A small fraction (less than 5%) of tpl embryos developing at 29°C arrest during embryogenesis. The arrested embryos express LENNY throughout a greater fraction of the embryo than do the non-arrested embryos (Fig. 6G,H). One interpretation of this arrested phenotype is that it reflects an early and more complete transformation of apical to basal fates and that such a transformation results in the arrest of embryonic growth. However, it is also possible that a second, unrelated mutation is responsible for the arrest or that the topless mutation interferes with the viability of early embryos in a way that is unrelated to the specification of apical or basal fates.

The J1092 marker is expressed in the embryo in presumptive root cap cells (Fig. 6C) (described by J. Hasseleff, http://www.plantsci.cam.ac.uk/Hasseleff/Home.html). Thirty seven percent (15/41) of homozygous topless mutant embryos developing at 29°C show expression of the J1092 reporter at both poles of the embryo by 168 hours postfertilization (Fig. 6J,K).

In summary, markers for basal root fates were often found expressed in the region of the developing apical root of topless mutants. Taken together with the experiments with apical

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**Fig. 5. Gene expression patterns in topless mutant embryos.**

(A) Expression of STM in wild-type heart stage embryo. (B) Expression of STM in a topless embryo (sibling to embryo in A). (C) UFO expression in topless mutant embryo. (D) UBIQUITIN expression in topless mutant embryo. (E) ANT expression in wild-type heart stage embryo. (F) ANT expression in topless mutant. (G) KNAT1 expression in wild-type walking stick stage embryo. (H) KNAT1 expression in topless embryo. (I) SCARECROW expression in wild-type globular embryos. (J) SCARECROW expression in topless mutant embryos. Scale bars: in A, 25 μm for A-F; in G, 50 μm for G,H; in I, 25 μm for LJ.
markers, we conclude that there is a transformation of the apical region of the embryo from a shoot ‘primordium’ into a root ‘primordium’ in topless mutant embryos.

**Expression of the DR5 auxin reporter in topless mutants**

A high auxin to cytokinin ratio will promote root development over shoot development in cultured cells (Skoog and Miller, 1957). One simple explanation for the mechanism of action of the topless mutation is that it increases the auxin concentration in apical regions of the embryo.

The activity of the DR5::GUS construct appears to reflect high levels of free auxin in the root (Sabatini et al., 1999). This synthetic promoter consists of seven tandem repeats of an element found in auxin responsive promoters (Ulmasov et al., 1997a). This sequence binds to a family of AUXIN RESPONSE FACTORS (ARF) thought to be involved in transducing the auxin signal. Sabatini et al. (Sabatini et al., 1999) have shown that this reporter is active in the presumptive root in the late heart stage embryo.

If the tpl-1 mutation acts to increase auxin levels in the apical region of the embryo and this, in turn, causes the apical pole to develop root fate, we might expect to see an early and broad accumulation of GUS activity in the apical regions of topless1:DR5::GUS embryos. To test this hypothesis, we allowed topless1:DR5::GUS embryos to develop at restrictive temperature (29°C) and assayed them for GUS expression at 168 hours post-fertilization. While all embryos expressed the reporter in the basal root, only two out of 100 embryos expressed GUS in the developing apical root of the topless mutant embryos at this timepoint. Recall that a much larger proportion of topless GUS in the developing apical root of the reporter in the basal root, only two out of 100 embryos expressed

**The topless-1 mutation partially suppresses the monopteros mutant phenotype**

The MONOPTEROS gene encodes a member of the ARF family of transcription factors and is thought to be responsible for transducing auxin signals crucial for root development (Hardtke and Berleth, 1998). monopteros mutants fail to develop a root and most of the hypocotyl during embryogenesis.

MONOPTEROS encodes a protein that has a DNA-binding domain at its N terminus, and a dimerization domain at its C terminus (Hardtke and Berleth, 1998). Between these two domains is a predicted activation domain. The rootless mutation (Barton and Poethig, 1993) is an EMS-induced allele of MONOPTEROS. PCR amplification of the MONOPTEROS-coding sequence in rootless mutants showed that these mutants carry a stop codon at position 594 of the polypeptide [Q594 (CAG) to STOP (TAG)]. This is downstream of the DNA-binding domain and upstream of the C-terminal dimerization domain. We have renamed this allele monopteros-7 (mp-7).

To determine if the topless-1 mutation acts through the monopteros pathway, we created mutants homozygous for both tpl-1 and mp-7 (Fig. 7). Self progeny from a plant of genotype tpl-1 mp-7/tpl-1 MP+ were scored. (Both genes are located on
chromosome 1.) At both 24 and 29°C, a quarter of the self-progeny lacked roots and were assumed to be the tpl-1 mp-7/tpl-1 mp-7 double mutants. Thus, the tpl-1 mutation did not restore the ability to make a basal root to monopteros mutants. By contrast, the tpl-1 mp-7 double mutants made apical roots similar to mp-7 single mutants (at 29°C). Thus, basal root formation is dependent on wild-type MONOPTEROS activity in topless mutants but apical root formation is not.

The tpl-1 mutation partially suppresses the hypocotyl defect seen in monopteros mutants: tpl-1 mp-7 double mutants were able to form more hypocotyl tissue than mp-7 single mutants (Fig. 7). Whereas only 9% (8/86) of monopteros single mutants made any detectable vasculature in the basal ‘stub’ typically found in mp mutants, 97% (102/105) of tpl-1 mp-7 mutants showed the presence of hypocotyl-associated vascular development (Fig. 7; this experiment was performed at 24°C). The partial suppression of the monopteros hypocotyl defect by tpl-1 shows that the tpl-1 mutation can have effects on basal portions of the developing embryo and that mutant tpl-1 activity can partially bypass the requirement for MONOPTEROS activity in the development of the hypocotyl and its vasculature.

Temperature shift experiments show that the tpl mutation can mediate its effects late in embryogenesis

The temperature sensitivity of the tpl-1 mutation allows for temperature shift experiments to determine at what point the tpl mutation affects the development of the apical pole of the embryo. For this experiment, homozygous tpl-1 plants were grown at 24°C. They were then pollinated and placed at 29°C or 17°C. Embryonic stages at the time of shift were determined by dissecting embryos from siliques pollinated and shifted at the same time as the experimental cohort.

Embryos allowed to develop at 17°C from fertilization onwards did not develop apical roots (Fig. 8). Embryos that experienced high temperature early in embryogenesis, but developed at 17°C from the 32-cell stage and onwards, developed a small fraction of double root seedlings. This indicates that tpl mutant activity, or synthesis of mutant TOPLESS product, during early stages of embryogenesis has the potential to affect development in an irreversible way. A majority of embryos shifted from 29°C to 17°C at the torpedo stage developed apical roots, indicating that the shift to permissive temperature was often not sufficient to restore normal shoot development after this point.

The experiment in which embryos were shifted from 17°C to 29°C at varying time points gave largely reciprocal results. Nearly all of the embryos that developed at 29°C from fertilization and on developed as double root embryos (Fig. 8). Shifting to 29°C during the 16-32 cell stage also resulted in a majority of double root seedlings, indicating that topless mutant activity or synthesis could alter fates at the globular stage. Surprisingly, shifting embryos from 17°C to 29°C as late as the transition to midheart stage could cause many of the embryos to make apical roots. This is surprising because transition and heart stage embryos express normal apical

Fig. 7. Phenotypes of mutants homozygous for both tpl-1 and mp-7. (A) Seedling to left is mp-7 homozygote. Seedling to right is mp-7 tpl-1 homozygote. Both seedlings developed at 24°C. The double mutant has a more well-developed hypocotyl than the mp-7 single mutant. (B) Cleared mp-7 single mutant (grown at 24°C) showing lack of vasculature in the hypocotyl ‘stub’. (C) Cleared mp-7 tpl-1 double mutant (grown at 24°C) showing vasculature extending down into the more well developed hypocotyl. (D) Homozygous mp-7 single mutant grown at 24°C. The double mutant has a more well-developed hypocotyl than the homozygote. Both seedlings developed at 24°C. The double tpl-1 mp-7 double mutant grown at 29°C. (E) Homozygous mp-7 tpl-1 double mutant grown at 29°C. The double mutant is able to develop an apical root. Scale bar: 200 μm in A; 100 μm in B,C; 50 μm in D,E.

Fig. 8. Effect of temperature shift on topless embryo development. Number of embryos that developed an apical root when shifted from 29°C to 17°C at fertilization, 0/126; at 8-16 cells, 6/127; at 16-32 cells, 8/154; at 32-64 cells, 4/126; at 64 cells to transition stage, 26/175; at transition to early heart stage, 43/103; at midheart stage, 48/91; at torpedo stage, 74/116. Number of embryos that developed an apical root when shifted from 17°C to 29°C at fertilization, 142/149; at 8-16 cells, 93/143; at 16-32 cells, 113/134; at 32-64 cells, 46/85; at 64 cells to transition stage, 92/195; at transition to early heart stage, 31/143; at midheart stage, 18/127.
the CUP-SHAPED COTYLEDON 1 and CUP-SHAPED COTYLEDON 2 genes; when simultaneously mutant, cue1 and cue2 cause the development of a cup-shaped cotyledon, failure to express STM and failure to form a SAM (Aida et al., 1997; Aida et al., 1999). This phenotype is seen in some tpl-1 mutants. It will be interesting to find out whether tpl-1 mutants are affected in CUC expression.

The effect of tpl-1 on the development of the hypocotyl is more subtle. With the exception of the double root seedlings, tpl-1 mutant hypocotyls appear superficially normal. However, developing topless embryos show greatly reduced KNAT1 transcript, a transcript normally specific to the developing hypocotyl. (Failure to express KNAT1 by tpl-1 mutant embryos is not a secondary consequence of failure to form a SAM, as sim2 mutants are able to express KNAT1 (J. L. and M. K. Barton, unpublished). Additional evidence for action of the tpl-1 mutation in the region of the developing hypocotyl is seen in the ability of the tpl-1 mutation to partially restore hypocotyl development in monopteros mutants.

Polarity in the embryo remains susceptible to tpl mutant activity until the heart stage

Temperature-shift experiments of developing topless embryos show that the probability of apical root development can be influenced by temperature even after cotyledons have begun to develop. Thus, a significant number (20%) of seedlings will develop apical roots when topless embryos are shifted to restrictive temperature at the transition to early heart stage, a stage when the presumptive cotyledons are beginning to grow out. This result strongly suggests that cells initially developing with apical fates are redirected into root fates in tpl embryos. Several other observations are consistent with this as well. First, ANT, a cotyledon marker, is expressed in all topless embryos even in conditions in which the embryos will develop apical roots. Second, the development of the apical root seems to occur later than that of the primary root in many topless embryos. This is indicated by our finding that despite the fact that a majority of topless embryos develop into double-root seedlings, a minority of topless embryos show expression of root specific molecular markers by 168 hours (7 days) post-fertilization. Finally, the tpl-1 apical root appears to derive from a larger group of cells than does a normal root. In the embryo, the tpl apical root is morphologically broader than a normal root and lacks the cellular organization of a normal root. Development from a larger group of cells is expected if the apical root develops from the more expanded apical region of a transition stage embryo.

Given the apparent respecification of shoot structures into root structures at the apical pole of at least some topless embryos, it will be interesting to learn whether the shoot to root transformation requires first dedifferentiation followed by redifferentiation, or whether apical cells switch fates directly, without undergoing an intervening stage of dedifferentiation.

One can speculate that the reason the double root embryos survive to be recognized at all is, in fact, because they undergo such a re-specification. If the apical pole of the globular embryo is mis-specified as ‘basal’ from the outset, the resulting ‘double-basal’ embryo may fail to develop sufficiently to complete embryogenesis. For example, failure to establish polarity along the apical/basal axis correctly might prevent growth of the embryo. The observation of arrested embryos

DISCUSSION

Homozygous topless mutant embryos exhibit transformation of shoot to root at the apical pole

The tpl-1 mutation causes dramatic changes in the development of the Arabidopsis embryo. The most dramatic and novel topless phenotype is seen at high temperature where embryos develop apical roots in place of shoot apical meristem and cotyledons. The apical root that develops has a recognizable root cap, exhibits normal gravitropism, has root hairs and shows generally normal morphology, although it may be wider than the wild-type root. In agreement with these gross characters, developing tpl-1 embryos show changes in pattern formation as detected with molecular markers for regional identities.

To our knowledge, this is the first example of a mutation that causes the transformation of apical into basal fates in the embryo of any plant species. The double root phenotype indicates that the tpl-1 mutation can bypass the requirement for any basal, root promoting, positional cues laid down in the egg cell or in the early zygote. Analogous mutations that cause transformation of basal into apical fates have not yet been discovered but the existence of the tpl-1 mutation suggests the existence of a master regulatory network that can switch relatively cleanly between apical and basal fates.

At lower temperatures, the seedlings are missing the shoot apical meristem and/or the cotyledons normally found at the apical pole. The SAM is more likely to be missing than are the cotyledons. Consistent with this, expression of the SAM markers STM and UFO was completely absent in all mutants, even at 24°C. Thus, tpl mutants are unable to specify SAM identity during embryogenesis. By contrast, topless mutant embryos appear to be able to specify cotyledon identity as ANT expression, although somewhat abnormal in pattern, is present. Surprisingly, ANT was even found to be present in embryos developing at 29°C, even though most of these embryos develop apical roots in place of cotyledons.

The abnormal pattern of ANT expression in topless mutant embryos, expression throughout the apical dome instead of only in a peripheral ring, indicates that pattern formation is defective not only in the apical-basal dimension but also in the radial dimension.

As loss of STM function is known to cause cotyledon fusion, the cotyledon defects may in part be a result of failure of homozygous tpl-1 seedlings to accumulate STM product. However, the severity of cotyledon fusion in tpl-1 seedlings can go well beyond what is seen in sim mutants, indicating that the regulation of other genes must be affected. Candidates include...
expressing root markers throughout the embryo is one line of evidence for such an effect. However, it will be crucial to determine that this arrest is due to the *tpl-1* mutation and not to a second unrelated mutation in this genetic background.

**Experiments fail to link the *topless* 1 mutation to alterations in auxin signaling**

It has long been recognized that a high auxin to cytokinin ratio will promote root development over shoot development in cultured cells (Skoog and Miller, 1957). One possibility is that *TPL* interferes with the perception of auxin to cytokinin ratios. *tpl* mutant embryos could have altered distribution patterns of these hormones, increased sensitivity to auxin or reduced sensitivity to cytokinins.

Our results do not support the simple hypothesis that the *topless*-1 mutation increases auxin levels in the apical pole of the embryo and thereby causes apical root development. If this were true, we would have expected to see early and ectopic expression of the DR5::GUS reporter; expression of this reporter is believed to reflect high auxin concentrations in the root (Sabatini et al., 1999). Instead, while a significant fraction of *topless* 1 embryos had begun to differentiate root fates by 168 hours postfertilization (as determined by the J1092 GFP and LENNY reporters), only a small fraction had begun to express the DR5::GUS promoter. One problem with the interpretation of this experiment is that the DR5::GUS promoter may not function as a reliable reporter for high free auxin levels in apical regions of the embryo.

The *MONOPTEROS* gene product has been implicated in auxin action in the developing root (Hardtke and Berleth, 1999). *MONOPTEROS* is required for the proper development of vasculature in the embryo and for the development of basal (hypocotyl and root) fates during embryogenesis. Many classical studies have indicated that auxin promotes both vascular and root development when exogenously applied. The finding that the *MONOPTEROS* gene product is a transcription factor that is similar to transcription factors believed to transduce auxin signals is further evidence that *MONOPTEROS* promotes vascular and root development through an auxin-regulated mechanism.

The *MONOPTEROS* gene product was not necessary for the formation of the apical root in *topless* mutant embryos. This is similar to the finding that *MONOPTEROS* is not required for ectopic root formation from the hypocotyl and cotyledons in *superroot* mutants (M. K. B., unpublished), the difference being that the apical root in *topless* mutants forms during embryogenesis, while the ectopic roots in *superroot* mutants form postembryonically. As there are many ARF-like genes in the *Arabidopsis* genome, it is possible that one or more homologous genes substitute for *MONOPTEROS* function when roots are formed from apical regions of the plant. Alternatively, *topless* may bypass the requirement for *ARF-LIKE* genes in apical root formation. By contrast, *topless* could not bypass the requirement for *MONOPTEROS* in the formation of the basal root.

*topless* monopteros double mutant individuals make more hypocotyl tissue than *monopteros* single mutants, indicating that the *topless* 1 mutation partially suppresses the monopteros phenotype. This could indicate that the *tpl-1* mutation activates a developmental pathway that can substitute in part for *MONOPTEROS* activity in hypocotyl development. Alternatively, the *topless* mutation may boost the activity of whatever residual *MONOPTEROS* activity exists in the *monopteros*–7 mutant.

In summary, the relationship of *topless* mutant activity to auxin signaling in root development remains unclear. Preliminary observations of *axr1 topless* double mutants show only additive interactions (J. L., unpublished), indicating that the two processes may indeed be independent of one another. Consistent with the notion of independent action of *topless* and auxin signaling, treatment of developing Indian mustard embryos with auxin transport inhibitors resulted in ball- or tube-shaped seedlings, but not in double rooted seedlings, as seen in *topless* mutants (Liu et al., 1993).

An additive action of *topless* and auxin signaling could also explain the temperature sensitive nature of the *tpl-1* mutation. Gray and co-workers have described auxin-dependent processes that are enhanced by high temperature (Gray et al., 1998). In addition, they measured higher levels of auxin in plants grown under high temperatures than under low temperatures. Thus, a combination of a mutant *tpl* gene product plus enhanced sensitivity to auxins and/or higher auxin levels at high temperatures might act in combination to explain the temperature sensitivity of the *topless* phenotype.

The role of **TOPLESS** in wild-type embryo development

The **TOPLESS** locus is currently represented by only one mutant allele. This allele is temperature sensitive and also shows a weak dominant effect. Additional screens for *topless*-like mutants have failed to turn up more alleles of the **TPL** locus, indicating that the change in *tpl-1* is likely to be a rare type of mutation (J. L. and M. K. B., unpublished).

Given the unique nature of this allele, it is difficult to determine the wild-type function of **TPL**. Several roles are possible, depending on the nature of the *tpl-1* mutation. If *tpl-1* is a loss-of-function or dominant negative mutation, the wild-type role of *tpl-1* is to confer shoot identity (or repress root identity) to the apical pole. However, if the mutation elevates wild-type **TOPLESS** activity or makes **TOPLESS** activity spatially or temporally constitutive, the wild-type role would be to confer root fate to the basal pole. Given the difficulty in obtaining new alleles, the molecular cloning of the **TOPLESS** locus is likely to be the most direct route to determining its wild-type function.

Rare dominant alleles of genes are difficult to obtain except through the use of efficient point mutagens such as EMS. Such alleles provide important clues to gene function that are easily missed in standard reverse genetic studies that rely on ‘knock-out’ or overexpression strategies. Examples of rare dominant mutations that disrupt regulation of protein products are those in the **ETR1** (ethylene resistant 1) and **PHABULOS**A genes (Bleecker et al., 1988; Chang et al., 1993; McConnell and Barton, 1998). Interestingly, both of these gene products appear to act as receptors. Whatever role **TOPLESS** plays in wild-type development, understanding and manipulating the *tpl-1* gene should lead to insights in embryonic pattern formation, as well as the ability to manipulate the generation of new root apical meristems from shoot tissue.

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