

Auxin and the Communication Between Plant Cells

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Abstract Multicellularity allows one to assign different functions to the individual cells. Cell fate could be defined by a stereotypic sequence of cell divisions or it might arise from intercellular communication between cells. Patterning in the totipotent plant cells results mainly from coordinative signals. Auxin is central in this respect, and this chapter ventures to give a survey on the role of auxin as a coordinative signal that regulates patterning of cell differentiation, cell division and cell expansion.

Abbreviations 2,4-D: 2,4-Dichlorophenoxyacetic acid; ARF: ADP-ribosylation factor; ARP: Actin-related protein; BFA: Brefeldin A; GFP: Green fluorescent protein; IAA: Indole-3-acetic acid; NAA: 1-Naphthaleneacetic acid; NPA: Naphthylphthalamic acid; RFP: Red fluorescent protein; TIBA: 2,3,5-Triiodobenzoic acid

1 Introduction 15

The polar flux of auxin has been used for more than 375 million years to generate and regulate the pattern of vascular differentiation of parenchymatic cells and thus coordinates the organization of the telomes, the building block of cormophytic land plants. In addition to the patterning of vasculature, auxin mediates the coordinative signalling that controls phyllotaxis, the formation of new leaves according to an orderly, species-dependent pattern. The phyllotactic pattern is shaped by competition of young primordia for free auxin, such that the neighbourhood of an existing primordium will be depleted of auxin. Since auxin limits the formation of new primordia, this simple mechanism ensures elegantly that new structures will be laid down at a minimal distance from preexisting primordia.

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26 Polar auxin transport can regulate the synchrony of cell divisions, with actin
27 organization emerging as a central factor defining the pattern of cell division, probably
28 by polarizing the flow of vesicles that deposit auxin-efflux carriers to the cell pole
29 and thus determining the directionality of auxin efflux. Since the organization of
30 actin, in turn, is regulated by auxin, a feedback loop is established that contains
31 auxin-efflux carriers, intracellular auxin and actin filaments as central elements.

32 Regulated cell expansion represents the central adaptive response of the sessile
33 plants to environmental challenges and is therefore highly responsive to stimuli,
34 such as light or gravity. These adaptive responses involve a spatiotemporal pattern
35 of cell expansion, which is most evident for tropistic curvature. Actually, auxin was
36 originally identified as a signal that coordinates the pattern of cell expansion.
37 The Cholodny–Went model explains tropism by a signal-induced redistribution of
38 auxin fluxes across the stimulated organ. Although the Cholodny–Went model is
39 repeatedly disputed mainly because of discrepancies between the observed response
40 (a gradient of growth) and the amplitude of the induced gradient of auxin, it is
41 shown that the model is still valid if the redistribution of auxin fluxes is complemented
42 by parallel gradients of auxin responsiveness.

43 The chapter ends with a speculative consideration of why, during evolution, such
44 a simple molecule as indole-3-acetic acid (IAA) has acquired such a central role for
45 intercellular coordination. This is attributed to the molecular properties of auxin that
46 determine its transport properties (multidirectional influx through an ion-trap mechanism,
47 but unidirectional efflux through the localized activity of auxin-efflux carriers).
48 On the intracellular level, this system is able to establish a clear cell polarity from
49 even minute and noisy directional cues. On the level of tissues, this system is ideally
50 suited to convey lateral inhibition between neighbouring cells. It was sufficient to
51 put the localization of the efflux transporter under the control of auxin itself to reach
52 a perfect reaction-diffusion system *in sensu* Turing (1952). Such systems are able to
53 generate clear outputs from even minute and noisy directional cues and provide a
54 robust mechanism to generate patterns of cell differentiation, cell division and cell
55 expansion under the special constraints of plant development, such as signal-dependent
56 morphogenesis and the lack of specialized and localized sensing organs.

57 Plant morphogenesis is not based on fixed hierarchies – there is no such a thing
58 as a “Great Chairman” that assigns differential developmental pathways to the
59 individual cells. Plant cells rather “negotiate” on their individual developmental
60 fates in a fairly “democratic” manner with hierarchies being created ad hoc by
61 mutual interactions. It seems that auxin has evolved as a central tool for this “cellular
62 democracy” characteristic for plant development.

63 2 Plant Development and Cell Communication

64 “Why do cells exist?” – with this question Philip Lintilhac (1999) starts his thoughtful
65 essay on the conceptual framework of cellularity. Multicellularity initially probably
66 evolved as a strategy to increase in size and thus escape the fate of being eaten.

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During growth, the volume of a cell (its “internum” *in sensu* Lintilhac) increases with the third power of the radius; its surface, though, increases only with the second power of the radius. When a cell grows, an increasing gap between consumption (by the “internum”) and subsistence (through the boundary with the “externum”) has to be bridged that will limit further expansion of the cell. Multicellularity allows an increase of the surface in relation to the volume – for the cell population *as an entity*. This made it possible for the cell to become bigger, again for the cell population *as an entity*. The selective advantage (not to be devoured by predators) paid off for each individual cell. However, the full potential of this achievement emerged only when the individual cells of the newborn organism began to assign different functions to individual members of the population. For the individual cell, differentiation represents a risky investment, because it implies that specific (Lintilhac coined the term “hypercellular”) tasks have to be upregulated at the cost of other “hypocellular” functions that are downregulated and therefore have to be compensated by corresponding hypercellular output from neighbouring cells. This culminates in a situation where the individual cells cannot survive outside the organismal context.

Differentiation therefore requires an intensive flow of information between individual cells to maintain the subtle balance between hypercellular and hypocellular functions. Although in some systems the differentiation of individual cells seems to follow a predetermined internal programme, cell–cell communication is important at least in the initial phase, when this programme is defined and triggered. Plant cells with their principal totipotency and their comparatively diffuse differentiation have to be especially communicative. Owing to their developmental flexibility, the balance between hypercellular and hypocellular functions has to be reestablished continuously. It thus seems that cell differentiation in plants resembles more or less the ancestral situation of multicellularity. In addition, plant cells are immobile, such that temporal patterns of differentiation become manifest morphologically and are not obscured by cell migrations.

The primordial form of cell differentiation is developmental dichotomy as characteristically observed during the first formative cell division of zygotes or spores in many algae, mosses and ferns or during the first division of the angiosperm zygote. In the Volvocales, a monophyletic clade of the green algae, it is still possible to follow the evolutionary line from a cell population over cell colonies (consisting of equivalent members that are completely autonomous) to a true organism, where two cell types are coupled by hypocellular and hypercellular interactions. Genetic analysis of differentiation mutants in *Volvox carteri* has uncovered a transcription factor, *regA*, repressing nuclear encoded genes of the chloroplast in mobile, somatic cells such that growth of these cells is suppressed, leading to a delayed cell cycle (Kirk 2003). In contrast, a group of four or five *late gonidia* factors suppress the motile phase in reproductive cells and thus promote their division. The activities of *regA* and *lag* differ as early as from the first division of the mature gonidium. This primary developmental dichotomy is under the control of two or three *gonidialess* factors – mutations in those genes render the first division symmetric such that the resulting daughter organism lacks reproductive cells. In fact, the dichotomy of

112 the first gonidial division is a cornerstone of August Weismann's concept of inheritance,
 113 where he defined the separation of the differentiating, but mortal *soma* from the
 114 non-differentiating, but immortal *germ lines* a primordial event of multicellular
 115 development (Weismann 1894).

116 Developmental dichotomy could be based on a gradient of developmental
 117 determinants within the progenitor cell that are then differentially partitioned to the
 118 daughter cells (formative cell division). According to this mechanism, the ultimate
 119 cause for differentiation would reside in cell lineage (Fig. 1a). Alternatively,
 120 developmental dichotomy could arise from communication between initially equipotent
 121 daughter cells and therefore would be independent of cell lineage (Fig. 1b).

122 As diverse as these two mechanisms might appear, it can be difficult to discriminate
 123 between them in nature since the commitment for a certain developmental pathway
 124 and the manifestation of this commitment as differentiation are not always clearly
 125 separated in time. However, the principal totipotency of plant cells is easier to reconcile
 126 with a model where differentiation is not defined a priori by a formative division
 127 (Fig. 1a), but a posteriori by intercellular communication (Fig. 1b).

128 The impact of intercellular communication on differentiation is heralded in the
 129 (prokaryotic, but plant-like) cyanobacteria during the differentiation of heterocysts.
 130 Heterocysts express (as hypercellular function) a nitrogenase that is able to release
 131 the constraints placed on cell division by the limited supply of bioavailable nitrogen.

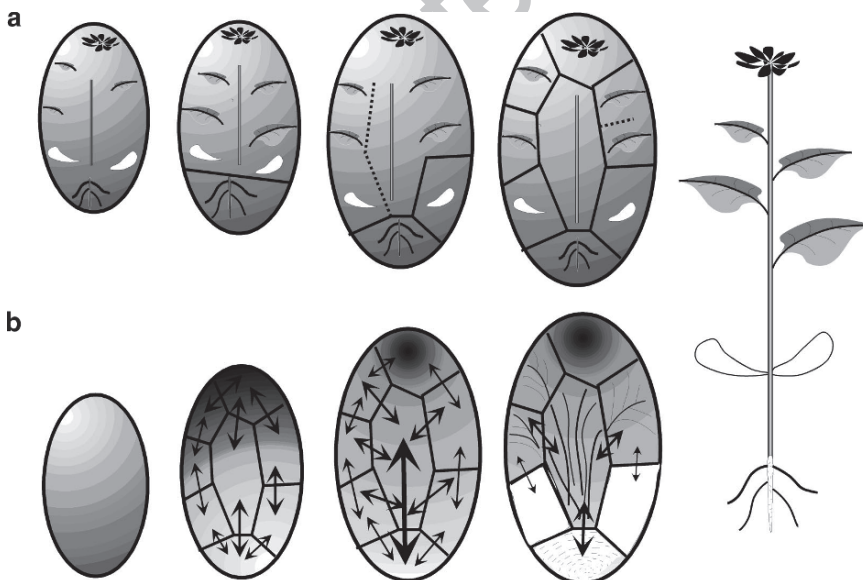


Fig. 1 Mechanisms for the establishment of developmental differentiation. (a) Mosaic development, where developmental fate is determined a priori and then assigned to individual daughter cells by a stereotypic sequence of formative cell divisions. (b) Regulative development, where developmental fate is not predetermined, but is defined a posteriori by communication between equipotent cells

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This nitrogenase dates back to the earliest, anoxic phases of life on this planet and is therefore highly sensitive to oxygen; therefore, to safeguard nitrogenase activity, any photosynthetic activity has to be excluded from heterocysts. These cells are therefore hypocellular with respect to assimilation. The balance between nitrogen export and assimilate import has to be maintained although the total number of cells grows continuously. This balance is kept by iterative patterning, whereby preexisting heterocysts suppress the differentiation of new heterocysts in a range of around ten cells. When, in consequence of cell divisions, the distance between them exceeds this threshold, a new heterocyst will differentiate between them. By the analysis of patterning mutants in *Anabaena* the factor responsible for this lateral inhibition could be identified as the diffusible peptide patS (Yoon and Golden 1998). Differentiation (including the synthesis of patS) will begin in clusters of neighbouring cells; however, one of these cells will advance and then suppress further differentiation in its neighbours (Yoon and Golden 2001). This demonstrates that the differentiation of a heterocyst is not predetermined, but is progressively defined by signalling between neighbouring cells.

Developmental dichotomy in the complete absence of a predefined gradient has been impressively demonstrated for the somatic embryogenesis of embryogenic carrot cell suspensions. Here a single cell can be induced to produce an entire embryo that is indistinguishable from a sexually produced plant. Similar to zygotic development, the initial event is an asymmetric cell division giving rise to a highly vacuolated basal and a smaller apical cell endowed with a very dense cytoplasm (McCabe et al. 1997). Whereas the vacuolated cell will undergo programmed cell death, this apical cell will undergo embryogenesis. The vacuolated cell expressed a surface marker that was recognized by the monoclonal antibody JIM8 that had originally been raised as a marker of cell fate in root development. By use of ferromagnetic antibody conjugates it was possible to remove cells expressing the JIM8 marker from the suspension. These cultures lost their embryogenic potential. However, a filtrate from a culture containing JIM8-positive cells was able to restore the embryogenic potential of the JIM8-depleted culture. The JIM8 marker, a small soluble arabinogalactan protein secreted by the vacuolated cells, was therefore necessary and sufficient to confer an embryogenic fate to the densely vacuolated apical cells. Thus, the formative division is controlled by intercellular signalling.

Whereas intercellular signalling is relatively evident in these two examples, it might be more widespread. The classic experiment to dissect the role of cell lineage versus intercellular signalling in animal embryology is to transplant tissue to a different site of the embryo and to test whether the explant develops according to position (favouring a signalling mechanism) or according to its origin, as would be expected for differentiation based on cell lineage (Spemann 1936). This experiment has rarely been undertaken in plants, and so intercellular signalling might have been overlooked in many cases. For instance, the highly stereotypic cell lineage in the root meristem of *Arabidopsis thaliana* seemed to indicate that here cell fate is defined by cell lineage that could be traced back to early embryogenesis (Scheres et al. 1994). However, by very elegant laser ablation experiments (Van den Berg et al. 1995) and the analysis of mutants with aberrant definition of tissue layers

177 et al. 2001) it could be shown that even in this case cell fate was defined by signals
 178 (such as the transcription factor *shortroot*) from adjacent cells.

179 Generally, the principal totipotency of plant cells is difficult to reconcile with a
 180 strong impact of cell lineage. Patterning in cells thus seems to result mainly from
 181 coordinative signals. However, as discussed in the next section, the impact of
 182 intercellular communication on development seems to reach beyond the realm
 183 of individual cells to the coordinative development of entire organs.

184 3 Auxin as a Pattern Generator in Cell 185 Differentiation I: Vasculature

186 As consequence of their light dependency, plants increase their surface in an
 187 outward direction, which means that they have to cope with a considerable degree
 188 of mechanic load. As long as they were aquatic, this was no special challenge,
 189 because mechanical strains were counterbalanced by buoyancy, allowing for
 190 considerable size even on the basis of a fairly simple architecture. However, the
 191 transition towards terrestrial habitats increased the selection pressure towards
 192 the development of flexible and simultaneously robust mechanical lattices. Plant
 193 evolution responded to this selective pressure by generation of load-bearing
 194 elements, the so-called telomes (Zimmermann 1965). These modules are organized
 195 around a lignified vascular bundle surrounded by parenchymatic tissue and an
 196 epidermis to limit transpiration (Fig. 2a). The telomes were originally dichotomously
 197 branched, but by asymmetric branching (“overtopping”) hierarchical branching
 198 systems emerged that were endowed with main and side axes. By planation and

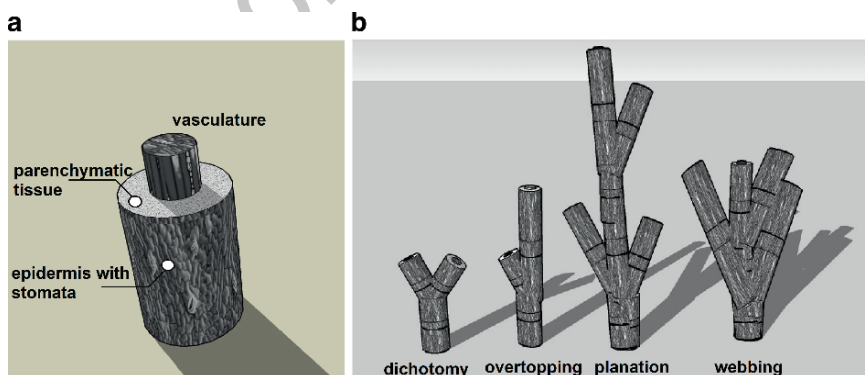


Fig. 2 Modular structure of terrestrial plants. The building block of cormophytes are the telomes (a), tubular elements organized around vasculature bundles that are surrounded by parenchymatic tissue and protected by an epidermis with stomata for the regulation of transpiration. By combination of the telomes in combination with simple modulations of their geometry (b) progressively complex hierarchical structures have been produced during the evolution of land plants



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subsequent fusion of the parenchymatic tissue (so-called webbing) the telomes developed into the first leaves (Fig. 2b). This is still evident in the leaves of certain ferns and the primitive gymnosperm *Ginkgo biloba*, where, occasionally atavistic forms are observed that uncover the original dichotomous telome structure. By spherical fusion and reduction of individual telomes globular structures arose that later evolved into sporangiophores and flower organs. Eventually, webbing of non-planar telomes generated the vasculature that has since been used throughout cormophyte evolution. In summary, the whole architecture of land plants derives from the patterned organization of these versatile modules. In other words, if one wants to understand the morphogenesis of land plants, one needs in the first place to understand the patterning of vessels as a core element of these building blocks.



Vessel patterning is central for the success of grafting in horticulture (Priestley Swingle 1929) and therefore shifted into the focus of botanical research many years ago. As long ago as the eighteenth century regenerative events in grafting were explained by a theory where two morphogenetic factors, a heavy “root sap” and a light “shoot sap”, moved towards the respective poles driven by gravity, accumulated there and triggered the formation of roots and shoots, respectively (Du Monceau 1764). In fact, the existence of such morphogenetic factors and their transport in the phloem was elegantly demonstrated by incision experiments (Hanstein 1860). By elaborate cutting and regeneration studies Goebel (1908) arrived at the conclusion that an apicobasal flux of an unknown substance defines the regeneration of new shoot and root elements. If this flux is interrupted or inverted, locally restricted inversion of shoot-root polarity becomes manifest as a gradient in the formation of vasculature and the ability to regenerate adventitious shoots or roots, respectively. The factor that was produced by developing leaves and that was able to induce the differentiation of new vasculature from parenchymatic tissue located basipetally of the leaf was later shown to be the transportable auxin IAA (Camus 1949). This finding opened up the possibility to experimentally manipulate the spatial pattern of vascular bundles, an approach that was exploited by Tsvi Sachs in a series of ingenious experiments. He could demonstrate that “differentiated vascular tissue whose source of auxin has been removed attracts newly induced vascular strands. This attraction is expressed in the joining of the new strands to the pre-existing vascular tissue. Differentiated vascular tissue which is well supplied with auxin inhibits rather than attracts the formation of new vascular strands in its vicinity” (Sachs 1968). This basic experiment and numerous

experimental derivatives culminated eventually in a canalization model of auxin-dependent patterning of vasculature: if, within an initially homogeneous distribution of auxin across the parenchymatic tissue, the polar flux of auxin is increased locally (for instance by blocking other drainage paths), the increase leads to accelerated differentiation of vessels at this site. Since those developing vessels can already transport more auxin per unit time, they will deplete the neighbouring areas of auxin. A few vessels will form and mutually compete for auxin. With time, the vessels differentiate progressively and, eventually, the pattern is stabilized by lignification (Sachs 1981).

244 The cellular basis of this drainage model is the polarity of vascular cells that are
245 aligned with the shoot–root axis. The vasculature of a leaf, however, does not reveal
246 such an obvious polarity and it was therefore not clear whether the auxin canalization
247 theory could be generalized to leaf veins as well. However, when transverse vascular
248 strands were investigated, they were found to include adjacent vessels with opposite
249 polarities that did not mature at the same time (Sachs 1975). Similar vessels without
250 clear polarity could be induced experimentally when the location of the auxin
251 source was changed repeatedly. Thus, the axis of a vessel seems to precede its
252 polarity, and the differentiation of a network without clear directionality (as typical
253 for dicotyledonous leaves) is thought to arise from non-synchronous auxin transport
254 across the leaf blade. Since auxin transport has been observed to oscillate in intensity
255 (Hertel and Flory 1968), the formation of an axial, but non-polar vessel could also
256 originate from auxin movement in opposite directions at different times even
257 through the same cells.


258 This model predicts that inhibition of polar auxin transport should impair the
259 differentiation of leaf veins. This has been tested experimentally in *Arabidopsis*
260 leaves treated with 1-naphthylphthalamic acid (NPA), an inhibitor of auxin
261 transport (Mattsson et al. 1999). When the concentration of the inhibitor was
262 raised, the vasculature was progressively confined to the leaf margin, indicating that
263 the central regions of the leaf were depleted of auxin. The canalization model is
264 further supported by a series of mathematical models that can explain a variety of
265 common venation pattern (for a recent review see Roeland et al. 2007). The molecular
266 base of canalization is generally seen by alignment of auxin-efflux transporters, [Au1]
267  as the PIN proteins, with the flux of auxin, such that these fluxes are amplified even
268  ner. In fact, PIN1 is polarized along the putative direction of auxin flow prior to
269 the formation of vasculature during early leaf development (Scarpella et al. 2006).

270 A central element of the canalization model is the feedback of auxin flux on cell
271 polarity. This implies that the cell responds to the flow rather than to the local
272 concentration of auxin, which poses sophisticated challenges for modelling.
273 Alternatively, PIN proteins could localize differentially to cell membranes depending
274 on the local auxin concentration in the cell adjacent to this membrane as proposed
275 for phyllotaxis (Jönsson et al. 2006). When this readout of local auxin concentration
276 is combined with an auxin-dependent expression of the channel, it is possible to
277 model channelling patterns that are consistent with those of the classical canalization
278 model (Roeland et al. 2007).

279  Although the cellular details of auxin channelling remain to be elucidated, it is
280  that this pattern generator is evolutionarily very ancient. Evidence for polar
281 auxin transport can be found in algae and mosses (for a review see Cooke et al. 2002)
282 and polar auxin transport has been proposed to be responsible for vascular differentia-
283 tion in early land plants (Stein 1993). In recent conifers or dicotyledonous plants
284 the vasculature follows straight lines, but forms characteristic whirlpools near buds,
285 branches or wounds when the presumed axial flow of auxin is interrupted. Identical
286 circular patterns also occur at the same positions in the secondary wood of the Upper
287 Devonian fossil progymnosperm *Archaeopteris*, thus providing the first clear fossil
288 evidence of polar auxin flow (Rothwell and Lev-Yadun 2005). Thus, already 375

million years ago ancient land plants used polar auxin flux as a tool to establish and 289
maintain a contiguous vascular pattern throughout their telomic modules. 290

4 Auxin as a Pattern Generator in Cell 291 Differentiation II: Phyllotaxis 292

 In addition to the patterning of vasculature, auxin is a central player in the coordina- 293
signalling that controls phyllotaxis, the formation of new leaves according to 294
an orderly, species-dependent pattern. It has been known for a long time that the 295
position of a prospective leaf primordium in the apical meristem is defined by 296
inhibitory fields from the older primordia proximal to the meristem (Schoute 1913). 297
This was demonstrated by isolation of the youngest primordium by tangential 298
incisions that shifted the position of the subsequent primordia (Snow and Snow 299
1931). At that time, this shift was interpreted in terms of a first available space 300
model, where the additional space created by the incision would allow the incipient 301
primordia to move to a position where they otherwise were excluded. However, this 302
result is consistent with inhibitory fields emanating from the primordia. There has 303
been a long debate on the nature of these inhibitory signals that were originally 304
thought to be chemical agents, but were later interpreted to be of mechanical nature. 305
Since a growing meristem is subjected to considerable tissue tension, the inhibition 306
could be merely mechanical, because the preexisting primordia would induce 307
stresses upon surrounding potential sites of primordium initiation. The expected 308
stress-strain patterns can perfectly predict the position of prospective primordia 309
(for a review see Green 1980). If the inhibition were mechanical, local release of 310
tissue tension by beads coated with extensin should alter phyllotaxis. In fact, such 311
beads could invert the phyllotactic pattern (Fleming et al. 1997). However, a closer 312
look showed that the extensin-induced structures did not always develop into true 313
leaves, but in some cases resembled mere agglomerations of tissue that did not 314
express leaf markers such as photosynthetic proteins. True leaf development was 315
only initiated when the extensin bead was placed in a site where according to the 316
natural phyllotaxis a primordium would have been laid down. This meant that mere 317
mechanical tension was not sufficient to explain phyllotaxis and this led to a 318
rehabilitation of chemical signals as the cause of the inhibitory field emanating 319
from preexisting primordia. Chemical inhibition was supported by studies in apices 320
that had been freed from primordia by application of auxin-transport inhibitors 321
(Reinhardt et al. 2000), an experimental system that allows study of the de novo 322
generation of a pattern without the influence of preexisting structures. In this system, 323
the coordinative signal was found to be auxin. However, against textbook knowledge, 324
the preexisting primordia did not act as sources, but as sinks for auxin. Within the apical 325
belt that is competent for the initiation of leaf primordia there is mutual competition 326
for auxin as a limiting factor and this competition is biased in favour of certain sites 327
(where, in consequence, a new primordium is initiated) by the preexisting primordia 328
that attract auxin fluxes from the meristem (Reinhardt et al. 2003). 329

330 The phyllotactic pattern could be explained by a mechanism where PIN1 that
 331 continuously cycles between an endocellular compartment and its site of activity at
 332 the plasma membrane acts as a sensor for intercellular auxin gradients (Roeland
 333 et al. 2007). When the endocytosis of PIN1 becomes suppressed by extracellular
 334 auxin (for instance through a membrane-bound or apoplastic auxin receptor; Fig.
 335 3a), auxin will be preferentially pumped upstream by an auxin gradient (Fig. 3b).
 336 In fact, the endocytosis of PIN1 has been shown to be suppressed by exogenous
 337 auxins (Paciorek et al. 2005) providing the positive amplification loop required for
 338 the auxin-dependent inhibitory field.
 339 Phyllotaxis and induction of vasculature are the two classic examples for
 340 auxin-dependent pattern formation. What can be generalized from these examples?
 341 Both patterns are highly robust against stochastic fluctuations of the input, they rely

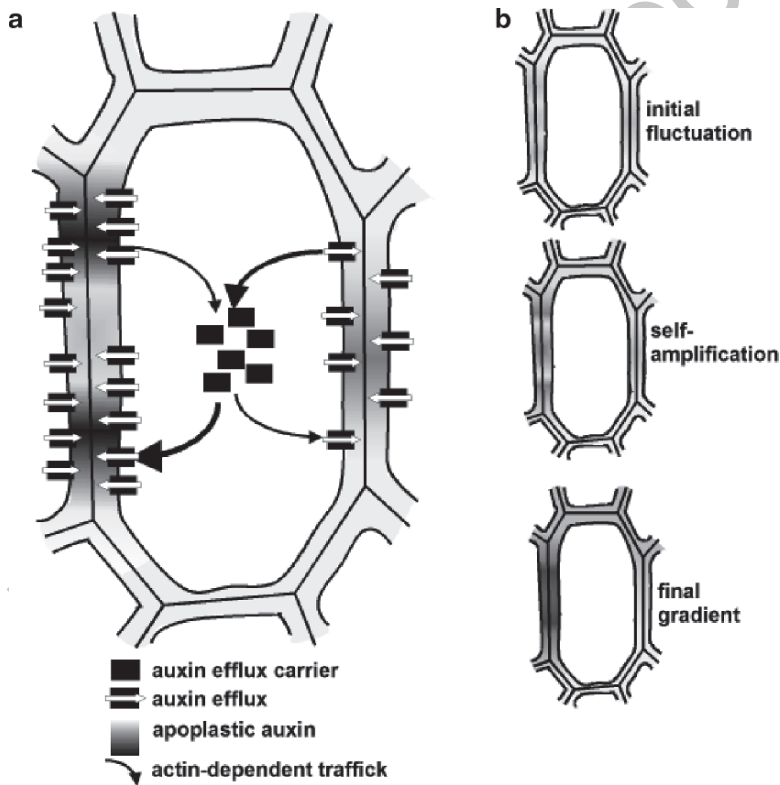


Fig. 3 Model for the self-amplification of transcellular auxin gradients. Auxin-efflux carriers cycle between the plasma membrane (their site of action) and an intracellular pool. Endocytosis of these carriers is locally inhibited by apoplastic auxin and is dependent on actin-mediated vesicle traffic (a). The competition between the two flanks of the cell for a limited number of the intracellular carriers in combination with local suppression of carrier endocytosis will amplify initial fluctuations of apoplastic auxin concentration progressively into clear gradients in the concentration of apoplastic auxin (b)

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on lateral inhibition between the patterned elements, and they culminate in qualitative decisions that are probably brought about by autocatalytic feedback loops. Such mechanisms can be described by the mathematics of reaction-diffusion systems that was adapted to biology (Turing 1952), and has been quite successfully used to model various biological patterns such as foot-head patterning in *Hydra* (Gierer et al. 1972), segmentation in *Drosophila* (Meinhard 1986) and leaf venation (Meinhard 1976). In these reaction-diffusion systems, a locally constrained, self-amplifying feedback loop of an activator is linked to a far-ranging mutual inhibition (Gierer and Meinhard 1972). Auxin-dependent patterning seems to follow this model, but differs in one aspect: rather than employing an actual inhibitor as a positive entity, in auxin-dependent patterning lateral inhibition is brought about by mutual competition for the activator.

5 Auxin as a Pattern Generator in Cell Division

In addition to cell expansion, auxin can induce cell division, a fact that is widely employed for tissue culture and the generation of transgenic plants. Investigation of lateral-root formation in *Arabidopsis* suggested that auxin regulates cell division through a G-protein-dependent pathway (Ullah et al. 2003, for a review see Chen et al. 2004). This was dissected further in tobacco suspension cells, early auxin signalling was dissected further, using the artificial auxins 1-naphthaleneacetic acid (NAA) and 2,4-dichlorophenoxyacetic acid (2,4-D). This study (Campanoni and Nick 2005) demonstrated that these two auxin species affected cell division and cell elongation differentially. NAA stimulated cell elongation at concentrations that were much lower than those required to stimulate cell division. In contrast, 2,4-D promoted cell division but not cell elongation. Pertussis toxin, a blocker of heterotrimeric G-proteins, inhibited the stimulation of cell division by 2,4-D but did not affect cell elongation. Conversely, aluminium tetrafluoride, an activator of the G-proteins, could induce cell division at NAA concentrations that were otherwise not permissive for division and even in the absence of any exogenous auxin. These data suggest that the G-protein-dependent pathway responsible for the auxin response of cell division is triggered by a different receptor than the pathway mediating auxin-induced cell expansion. The two receptors appear to differ in their affinity for different auxin species, with 2,4-D preferentially binding to the auxin receptor responsible for division and NAA preferentially binding to the auxin receptor inducing cell growth.

This bifurcation of auxin signalling (Fig. 4) appears to imply a differential interaction with the cytoskeleton as suggested by a recent detailed study on the effect of auxin on root growth in *Arabidopsis thaliana* (Rahman et al. 2007). When the contributions of cell division and cell elongation were assessed separately, the natural auxin IAA along with NAA and the auxin-transport inhibitor 2,3,5-triiodobenzoic acid (TIBA) were observed to inhibit cell elongation while leaving filamentous actin basically unaltered. In contrast, 2,4-D and the polar transport inhibitor NP inhibited cell division and at the same time eliminated actin filaments.

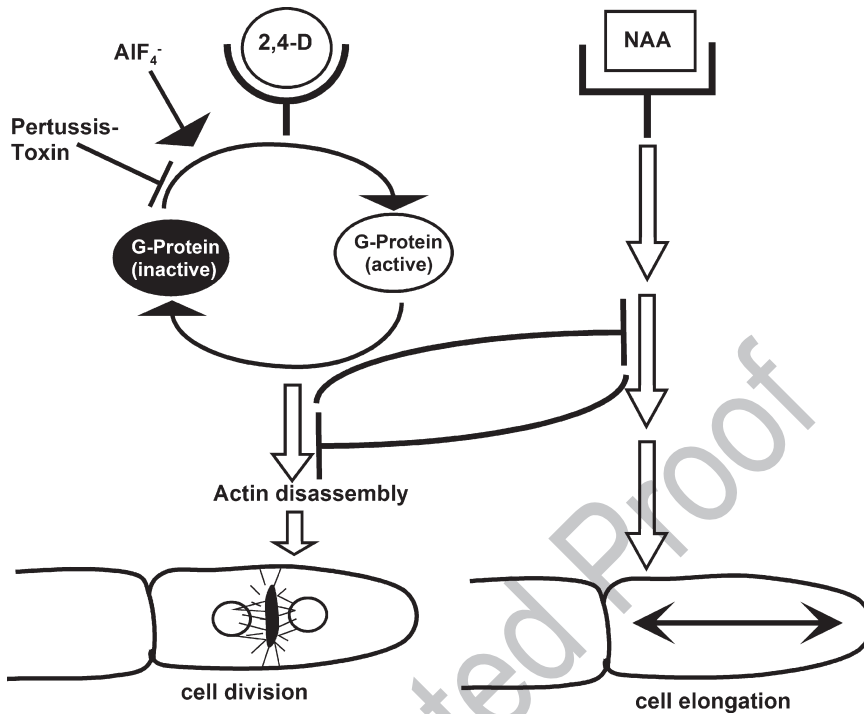


Fig. 4 Model for the bifurcation of auxin signalling in the regulation of cell division and cell elongation in tobacco cells according to Campanoni and Nick (2005) modified according to Rahman et al. (2007). The auxin receptor with high affinity for 1-naphthaleneacetic acid regulates cell elongation and is independent of G-protein activity and does not cause a disassembly of actin, whereas the auxin receptor with high affinity for 2,4-dichlorophenoxyacetic acid triggers a signal chain that involves the activity of a G-protein and triggers the disassembly of actin filaments. This signal chain is inhibited by pertussis toxin and is activated by aluminium tetrafluoride. Both pathways are mutually inhibitory

383 The root represents a very complex system consisting of different tissue layers
 384 that differ with respect to molecular machinery, auxin sensitivity and cytoskeletal
 385 organization. Moreover, the frequency of cycling cells, even in a rapidly growing
 386 root, is relatively modest, which makes it difficult to study the control exerted by
 387 intercellular auxin signalling on cell division on a quantitative level. However, a
 388 clear pattern of cell divisions is evident, with the cells of the quiescent centre acting
 389 as stem cells for the generation of proliferative tissues. As pointed out already, in
 390 the primary root of *Arabidopsis*, where this phenomenon has been dissected most
 391 intensively, this pattern can be traced back to early embryogenesis, whereas it
 392 seems to be more flexible in meristems of the Graminea. Nevertheless, the pattern
 393 of cell division is already established when the root meristem becomes accessible
 394 to cell-biological inspection and it is very difficult, if not impossible, to manipulate
 395 these patterns in a fundamental manner. Thus, root meristems represent a beautiful

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system to study pattern perpetuation, but for the analysis of pattern induction simpler systems that are less determined might be more appropriate. Suspension lines of tobacco are such models to study the primordial stages of division patterning and, in general, cellular aspects of cell division. These lines usually proceed from unicellular stages through a series of axial cell divisions towards cell files that are endowed with a clear axis and, in most cases, with a clear polarity. As will be explored in more detail below, these cell files are not a mere aggregation of autonomous, independent, cells, but display holistic properties such as an overall directionality and a pattern of cell division. In other words, these files are nothing other than a very reduced, but entire version of a multicellular “organism”. Owing to this extreme reduction in the level of complexity, it may be easier to study the intercellular negotiations of hypercellular and hypocellular functions rather than in a highly complex and differentiated meristem. Two tobacco cell lines have been studied in more detail with respect to cell–cell communication:

☰ The cell line VBI-0 (Opatrný and Opatrná 1976; Petrášek et al. 1998) derives from stem pith parenchyma, i.e. the cells that can differentiate into vascular tissue in response to auxin flow. These cells have preserved the ability to generate the structured cell-wall thickenings characteristic for xylogenesis (Nick et al. 2000). In the same way as its parenchymatic ancestor cells, this cell line grows in files where fundamental characteristics of patterning, such as clear axis and polarity of cell division and growth, are manifest. The progression into the culture cycle, the duration of the lag phase, the rate of cell division and the length of the exponential phase (Campanoni et al. 2003), but also cell polarity and axially (Petrášek et al. 2002), can be controlled by auxin. The cell files are formed from singular cells, such that positional information inherited from the mother tissue probably does not play a role. If there are patterns of competence within a cell file, they must originate de novo during the culture cycle.

☰ The widely used cell line BY-2 (Nagata et al. 1992) has generated a wealth of data on the role of phytohormones during the plant cell cycle. Compared with VBI-0, the temporal separation between cell division and cell expansion phases is less pronounced (probably as a consequence of the extremely high mitotic activity and short culture cycle). Moreover, the subsequent differentiation of these cells cannot be observed because they very rapidly lose viability if they are not subcultured directly after the logarithmic phase. However, BY-2 is transformed much more easily than VBI-0, such that a broad panel of different transgenic lines expressing fluorescently tagged marker proteins has become available. Moreover, although not as clearly manifest as in VBI-0, the basic features of patterning as well as file axis and polarity can be observed as well in this line.

During the work with these two cell lines, the cell divisions within the file were found to be partially synchronized, leading to a much higher frequency of cell files with even cell numbers than cell files with uneven cell numbers (Campanoni et al. 2003; Maisch and Nick 2007). The experimental data could be simulated using a mathematical model derived from non-linear dynamics, where elementary oscillators (cycling cells) were weakly coupled, and where the number of these oscillators was not conserved, but increased over time. The model predicted several non-intuitive

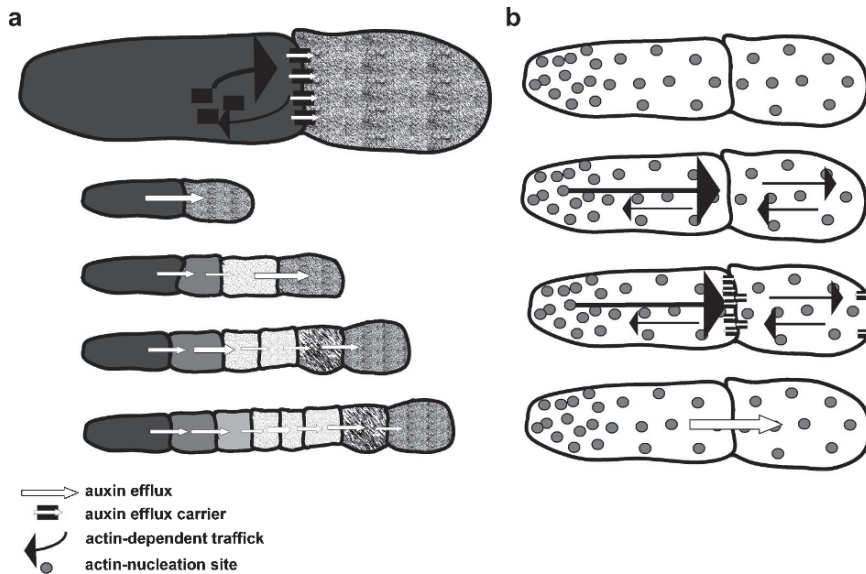


Fig. 5 Model for the regulation of cell division patterns in tobacco cell cultures by polar auxin transport. **a** Actin-dependent cycling of auxin-efflux carriers results in a polar distribution of the carrier and a polar flow of auxin through the cell file. Divisions of neighbouring cells are synchronized by this flow such that even cell numbers become more frequent than uneven cell numbers. **b** Actin-related protein 3 as marker for actin-nucleation sites is distributed in a gradient in the polarized tip cells, but not in the other cells of the file. The gradient of actin nucleation should result in a gradient of actin-dependent traffic that in turn will generate a graded distribution of auxin-efflux carriers such that auxin flow is polarized along the file axis

441 properties of the experimental system, among them that this coupling is unidirectional,
 442 i.e. that the coordinating signal was transported in a polar fashion. The coupling
 443 corresponds to a phase shift in the cell cycle, i.e. a dividing cell will cause its
 444 downstream neighbour to accelerate its cell cycle such that it will also initiate
 445 mitosis. The synchrony of cell divisions could be inhibited by low concentrations
 446 of the auxin-efflux inhibitor NPA. Although it has been known for a while that
 447 auxin is necessary for the progress of the cell cycle, and thus can be used to synchronize
 448 the cell cycle in plant cell cultures (for a review see Stals and Inzé 2001), this was
 449 the first time that auxin was shown to coordinate the divisions of adjacent cells.
 450 The modelling and the time courses of cell division showed that the noise in this
 451 system was considerable, with high variation in the cycling period over the cell
 452 population. Nevertheless, the division of adjacent cells was synchronized to such a
 453 degree that files with uneven cell numbers were rare compared with files with even
 454 numbers (Fig. 5a). Frequency distributions over the cell number per file thus exhibited
 455 oscillatory behaviour with characteristic peaks at the even numbers.

456 Since auxin efflux carriers cycle between the plasma membrane and an endocytotic
 457 compartment, auxin signalling has been linked to the organization of actin (for a
 458 review see Xu and Scheres 2005). However, this presumed link has recently been

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questioned by experiments, where PIN1 and PIN2 maintained their polar localization, 459
 although actin filaments had been eliminated by 2,4-D or NPA (Rahman et al. 2007). 460
 For the phytohormones TIBA and 2-(1-pyrenoyl) benzoic acid, it was shown very 461
 recently that they induce actin bundling not only in plants, but also in mammalian and 462
 yeast cells, i.e. in cells that are not to be expected to utilize auxin as a signalling 463
 compound (Dhonukshe et al. 2008). This has been interpreted as supportive evidence 464
 for a role of actin filaments in polar auxin transport. However, it was mentioned in the 465
 same work that NPA failed to cause actin bundling in non-plant cells, suggesting that 466
 its mode of action must be different. 467

Irrespective of the suggested direct effect of TIBA and 2-(1-pyrenoyl) benzoic acid 468
 on microfilaments, actin organization has been found to be highly responsive to 469
 changes in the cellular content of auxins (which would explain the NPA effect, for 470
 instance). This finding is actually quite old. During the classical period of auxin 471
 research, Sweeney and Thimann (1937) proposed that auxin might induce coleoptile 472
 growth by stimulating cytoplasmic streaming that is indeed very prominent in the 473
 coleoptile epidermis. In a series of publications, the late Kenneth Thimann returned 474
 to this idea and showed that elimination of actin very efficiently blocked auxin-dependent 475
 growth and argued that microfilaments are necessary for cell growth (Thimann et al. 476
 1992; Thimann and Biradivolu 1994). These findings contrasted with laser tweezer 477
 measurements, where the rigour of actin limiting cell expansion was shown to be 478
 released by auxin (Grabski and Schindler 1996). In the framework of this actin-rigour 479
 model, the elimination of actin would be expected to stimulate rather than inhibit 480
 auxin-dependent growth. On the other hand, at that time there was no alternative 481
 model that could explain how actin filaments would support cell growth. 482

To get insight into the role of actin in the control of cell growth, the phytochrome- 483
 triggered cell elongation of maize coleoptiles was studied in more detail (Waller and 484
 Nick 1997), leading to a physiological definition of two microfilament populations 485
 that were functionally different. In cells that underwent rapid elongation, actin was 486
 organized into fine strands that became bundled in response to conditions that inhibited 487
 growth. This transition was rapid and preceded the changes in growth rate. Moreover, 488
 this response was confined to the epidermis, i.e. to the target tissue for the signal 489
 control of growth (Kutschera et al. 1987). Later, these two actin populations could be 490
 separated biochemically owing to differences in sedimentability (Waller et al. 2002). 491
 The fine actin filaments correlated with a cytosolic fraction of actin, whereas actin 492
 became trapped on the endomembrane system and was partitioned into the microsomal 493
 fraction in conditions that induced bundling. The transition between the two states of 494
 actin could be induced, in a dose-dependent manner, by light (perceived by 495
 phytochrome), by fluctuations of auxin content, or by the secretion inhibitor brefeldin 496
 A (BFA). The bundling of actin was accompanied by a shift of the dose-response of 497
 auxin-dependent cell elongation towards higher concentrations and thus to a reduced 498
 auxin sensitivity *in sensu strictu*. This led to a model whereby auxin signalling caused 499
 a dissociation of actin bundles into finer filaments that were more efficient with 500
 respect to the polar transport of auxin-signalling/transport components. Thus, any 501
 modulation of cellular auxin content (such as that induced by phytohormones) is 502
 expected to repartition the ratio between bundled and detached actin filaments. 503

504 This short excursion makes clear that although the organization of actin seems
 505 to play a role in the polarity of auxin fluxes, there is also a clear effect of auxin on
 506 the organization of actin filaments. This bidirectionality in the relation between
 507 actin and auxin has to be considered to avoid flaws in the interpretation of inhibitor
 508 effects. The feedback circuit between auxin and actin was addressed using patterning
 509 of cell division as a sensitive trait to monitor changes of polar auxin fluxes (Maisch
 510 and Nick 2007). If actin were part of an auxin-driven feedback loop, it should be
 511 possible to manipulate auxin-dependent patterning through manipulation of actin
 512 (Fig. 5a). This prediction was tested using a transgenic BY-2 cell line stably
 513 expressing a fusion between the yellow fluorescent protein and the actin-binding
 514 domain of mouse talin (Ketelaar et al. 2004). In this cell line, the microfilaments
 515 were constitutively bundled, and the synchrony of cell division was impaired in
 516 such a way that the characteristic oscillations described above disappeared. When
 517 transportable auxin was added (auxin per se was not sufficient), both a normal
 518 organization of actin and the synchrony of cell division could be restored. This
 519 demonstrated that actin is not only responsive to changes in the cellular content of
 520 auxin, but that it also actively participates in the establishment of the polarity that
 521 drives auxin transport.

522 When actin organization is relevant for the synchrony of cell division (mediated
 523 by a polar transport of auxin), the factors that regulate the organization and polarity
 524 of actin filaments are highly relevant for patterning. A central player might be the
 525 actin-related protein (ARP) 2/3 complex, a modulator of the actin cytoskeleton shown
 526 by immunofluorescence to mark sites of actin nucleation in tobacco BY-2 cells
 527 (Fišerová et al. 2006). ARP2/3 caps the pointed end such that the actin filament
 528 grows in the direction of the barbed end. Tobacco Arp3 was cloned and fused to red
 529 fluorescent protein (RFP) as a marker for bona fide sites of actin nucleation (Maisch
 530 J, Fišerová J, Fischer L, Nick P, submitted). By biolistic transient transformation of
 531 tobacco cells it was possible, for the first time, to visualize ARP3 in living plant cells.
 532 With use of dual-fluorescence visualization of actin [by a green fluorescent protein
 533 (GFP) fusion of the actin-binding site of fimbrin] the RFP-ARP3 could be shown to
 534 decorate actin filaments in vivo. When actin filaments were transiently eliminated
 535 (either by treatment with cytochalasin D or by cold treatment) and then allowed to
 536 recover, RFP-ARP3 marked the sites from which the new filaments emanated.

537 With use of this marker, the behaviour of actin-nucleation sites could be
 538 followed through patterned cell division in comparison with AtPIN1::GFP-PIN1 as a
 539 marker for cell polarity. This uncovered a qualitative difference between the terminal
 540 (polarized) cells of a file and the (isodiametric) cells in the centre of a file (Fig. 5b).
 541 The density of ARP3 was increased in the apex of terminal cells in a gradient
 542 opposed to the polarity monitored by PIN1 (which was concentrated at the opposite,
 543 proximal cross wall). Upon disintegration of the file into single cells, the graded dis-
 544 tribution of ARP3 persisted, whereas PIN1 was redistributed uniformly over the plasma
 545 membrane of these cells. In contrast, the isodiametric cells in the file centre did not
 546 exhibit a graded distribution of the ARP3 signal, and the accumulation of PIN1 at the
 547 cross wall was much fainter than at the terminal cells, indicating that they are caused
 548 by residual amounts of PIN1 laid down by the (polar) progenitors of these cells.

[Au2]

[Au3]

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The relationship between actin, vesicle flow and polar auxin transport appears to be interwoven by a bifurcated signal chain: vesicle trafficking mediated by ADP-ribosylation factors (ARFs) is required for the polar localization of Rho-related GTPases in plants which control regulators of the ARP2/3 complex (Frank et al. 2004). On the other hand, ARF-mediated vesicle trafficking also controls the localization of PIN proteins which is known to rely on the activity of the serine-threonine kinase PINOID (Friml et al. 2004) and on the function of P-glycoproteins/multiple drug resistance proteins (Noh et al. 2001). When the function of these ARFs is impaired, in consequence of either treatment with the fungal toxin BFA or a mutation in one of the guanine nucleotide exchange factors that activate the ARFs, PIN1 becomes mislocalized and is trapped in intracellular compartments (Geldner et al. 2001). This cellular effect accounts for the phenotype of the corresponding *Arabidopsis* mutant, *gnom*, that suffers from a drastic loss of cell and organ polarity and, in consequence, is not able to establish an organized *Bauplan*. Thus, ARF-dependent vesicle flow controls actin nucleation (through the activity of the ARP2/3 complex) and, in parallel, the localization of PIN proteins. However, the initial cue that controls the spatial pattern of ARF activity remains unknown. ARP3 maintained an intracellular gradient in the polar terminal cells of BY-2, whereas PIN1 was redistributed (Maisch J, Fišerová J, Fischer L, Nick P, submitted), indicating that actin nucleation might be upstream of the events that culminate in a polar distribution of PIN1. However, owing to the split signalling of the ARFs on the Rho-related GTPases and on the ARP2/3 complex, ARP3 and PIN1 might as well be parallel downstream targets of unknown factors that are expressed in response to cell polarity.

[Au4]

Irrespective of these uncertainties in the molecular details, actin filaments have emerged as central players for the directional vesicle flow by which the polar localization of auxin-efflux carriers is established and perpetuated. The cycling of PIN1 is suppressed by exogenous auxin such that PIN1 remains longer in the plasma membrane (Paciorek et al. 2005) and is therefore able to pump auxin more efficiently into the apoplast. On the other hand, the localization of PIN1 depends on the activity of actomyosin and the organization of the actin tracks is in turn under the control of auxin. These interactions will therefore establish a feedback loop with auxin-efflux carriers, intracellular auxin and actin filaments as central elements (Fig. 3a). This feedback loop is nothing other than a reaction-diffusion system *in sensu* Turing and might represent the cellular pacemaker of auxin-mediated pattern formation.

6 Auxin as a Pattern Generator in Cell Expansion



Once a plant cell has been born by cell division, it undergoes rapid expansion by uptake of water. This expansion is impressive: plant cells can increase in size by up to 4 orders of magnitude (Cosgrove 1987). Regulated cell expansion represents the central adaptive response of the sessile plants to environmental challenges and is therefore highly responsive to stimuli, such as light or gravity, and internal factors, including developmental signals and plant hormones. Whereas the mechanisms

590 driving and regulating cellular expansion have been investigated in great detail over
 591 several decades, relatively little attention has been paid to the coordinative aspects
 592 of cell expansion. However, historically it was exactly this coordination of cell
 593 expansion that led to the discovery of auxin. In their famous *The Power of Movement*
 594 *in Plants*, the Darwins demonstrated for the phototropism of graminean seedlings
 595 that the direction of light is perceived in the very tip of the coleoptile, whereas the
 596 growth response to this directional stimulus occurs at the coleoptile base (Darwin
 597 and Darwin 1880). The signal transported from the tip to the base of the coleoptile [Au5]
 598 must transmit not only information about the fact that the coleoptile tip has perceived
 599 light, but also information about the direction of the light stimulus.

600 Simultaneously, but independently, Cholodny (1927), for gravitropism, and Went
 601 (1926), for phototropism, discovered that this transmitted signal must be a hormone.
 602 By means of the famous *Avena* biotest this hormone was later found to be IAA (Kögl
 603 et al. 1934; Thimann 1935). The Cholodny–Went model explains tropistic curvature
 604 by an alignment of auxin transport with the stimulation vector. The resulting gradient of
 605 auxin between the two flanks of the stimulated organ will then induce a growth
 606 differential that drives bending in the direction of the inductive stimulus.

607 Since its beginnings, the Cholodny–Wentmodel has been challenged by attempts
 608 to explain tropism independently of cell communication by mere summation of cell-
 609 autonomous responses (Fig. 6a). For instance, when light causes an inhibition of
 610 growth, a gradient of light should produce a gradient of growth that would not
 611 require the exchange of intercellular signals (Blaauw 1915). Alternatively, each cell
 612 could perceive the direction of the stimulus and produce a directionality on its own
 613 – without interaction with the other cells – and the individual cell polarities would
 614 then add up to the polarity of the entire organ (Heilbronn 1917). This debate stimulated
 615 an ingenious experiment by Johannes Buder, where the gradient of light across
 616 the tissue and the direction of light were opposite (Fig. 6b). He irradiated the
 617 coleoptile from inside-out using a prototype of a light-pipe (Buder 1920). Under
 618 these conditions, the coleoptiles bent towards the lighted flank, i.e. according to the
 619 gradient of light and opposite to its direction. The outcome of this experiment
 620 demonstrated clearly that the direction of light is sensed in the coleoptile tip owing
 621 to extensive communication between the perceptive cells and strongly argues against
 622 cell-autonomous models of tropistic perception (Nick and Furuya 1996).

623 The transverse polarity built up in response to phototropic or gravitropic stimulation
 624 in the perceptive tissue subsequently redistributes the basipetal flow of auxin and thus
 625 transmits the directional information into the responsive tissue at the coleoptile base.
 626 This gradient of auxin flow is well established, starting from bioassays (for instance
 627 Dolk 1936) and ending up with tracer experiments using radioactively labelled auxin
 628 (Goldsmith and Wilkins 1964; Parker and Briggs 1990; Iino 1991; Godbolé et al.
 629 2000) or direct measurements of free auxin across tropistically stimulated coleoptiles
 630 (Philippart et al. 1999; Gutjahr et al. 2005).

631 The Cholodny–Went model has been under continuous debate (see also [Au6]
 632 Trewavas 1992), mainly because there is a discrepancy in amplitude between the
 633 gradient of the growth rate and the gradient of auxin concentration. The difference
 634 in auxin content between the two flanks of a tropistically stimulated coleoptile is in the

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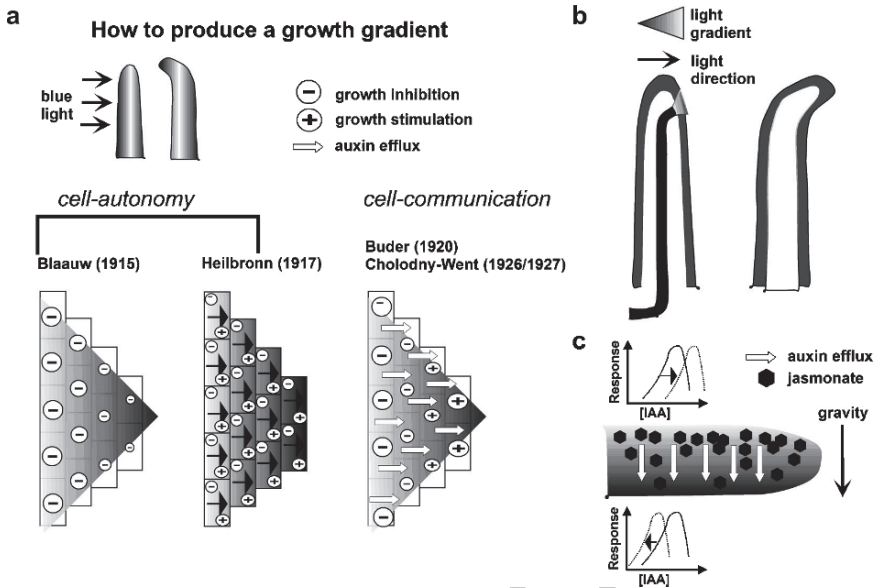


Fig. 6 Patterns of cell expansion during tropistic curvature of graminean coleoptiles. **a** Models for the formation of a growth gradient. According to Blaauw (1915), phototropic curvature emerges from a summation of growth inhibitions in response to the local intensity of light without interaction between cells. According to Heilbronn (1917), individual cells perceive the direction of light and respond by an intracellular gradient of growth. In contrast to these models that are based on complete cell autonomy, Buder (1920) explains curvature by interactions between individual cells across the coleoptile, and Cholodny (1927) and Went (1926) imply a lateral transport of a growth substance ("auxin"). **b** The experiment of Buder (1920), where the light direction and the light gradient across the organ are opposed. The bending is determined by the gradient, not by the direction of the light, contradicting the model postulated by Heilbronn (1917). **c** Extended Cholodny-Went model of gravitropism (according to Gutjahr et al. 2005). The lateral transport of auxin across the stimulated coleoptile is accompanied by a counterdirected gradient of jasmonate abundance and a gradient of auxin responsiveness across the tissue, with elevated responsiveness in the lower flank and reduced responsiveness in the upper flank

range of about 1:2 (Goldsmith and Wilkins 1964; Parker and Briggs 1990; Gutjahr et al. 2005), whereas growth is completely shifted from one flank to the other, i.e. the decrease in growth rate in one flank corresponds to the increase in growth rate in the other flank (Digby and Firn 1976; Iino and Briggs 1984; Himmelspach and Nick 2001). Elongation growth in coleoptiles increases more or less proportionally to the logarithm of auxin concentration (Wang and Nick 1998), such that the observed doubling of auxin concentration in the one flank would not succeed in causing the observed changes in growth rate. Moreover, when gravitropically stimulated hypocotyls (Rorabaugh and Salisbury 1989) or coleoptiles (Edelmann 2001; Gutjahr et al. 2005) were submersed in high concentrations of auxin, they showed positive gravitropism, i.e. they behaved as if they were roots. This is difficult to reconcile with a gradient of auxin as a unique cause for tropistic curvature.

647 With use of a classic biotest for auxin, the split-pea assay, in gravitropically
648 stimulated rice coleoptiles, it could be demonstrated that, in parallel to the
649 redistribution of auxin itself, a gradient of auxin responsiveness develops (Gutjahr
650 et al. 2005) with elevated responsiveness at the lower flank and reduced responsiveness
651 at the upper flank (Fig. 6c). This gradient of responsiveness can account for the
652 strong redistribution of growth even for relatively modest gradients of auxin. It can
653 even explain the peculiar sign reversal of bending for incubation with high concen-
654 trations of auxin beyond the optimum – for such superoptimal concentrations, the
655 elevated auxin responsiveness at the lower flank should result in an inhibition of
656 growth that is less pronounced in the upper flank, where the responsiveness is
657 lower. In parallel to the gradient of auxin, a counterdirected gradient of jasmonate
658 developed with higher concentrations at the upper flank as compared with the lower
659 flank. Jasmonate acts as a negative regulator for auxin responsiveness, because both
660 signal pathways compete for signalling factors such as AXR1 (Schwechheimer
661 et al. 2001). Thus, the observed jasmonate gradient might well account for the
662 observed gradient in auxin responsiveness across a gravitropically stimulated
663 coleoptile. To test this assumption, the jasmonate gradient was either equalized by
664 flooding the coleoptiles with exogenous methyl jasmonate or eliminated in
665 consequence of a mutation that blocks jasmonate synthesis (Gutjahr et al. 2005).
666 In both cases, the gravitropic response was delayed by about 1 h, but was eventually
667 initiated and proceeded normally. This indicates that the jasmonate gradient, although
668 not necessary for gravitropism, acts as a positive modulator. When auxin transport
669 was inhibited by NPA, the jasmonate gradient nevertheless developed, suggesting
670 that it is induced by gravitropic stimulation in parallel to and not in consequence of
671 lateral auxin transport.

672 In summary, the Cholodny–Went model has to be extended by signal-triggered,
673 modulative gradients of auxin responsiveness, but remains valid in its central statements.
674 This means that tropistic responses, representing nothing other than a patterned
675 distribution of cell expansion over the cross-section of the stimulated organ, can be
676 explained in terms of auxin-dependent cell communication.

677 The analysis of auxin-dependent cell communication in cell division has identified
678 a feedback loop between actin and auxin. This loop represents also a central element
679 of patterned cell expansion. Actually, it was cell elongation in coleoptiles where the
680 regulation of actin organization by auxin was discovered first (Sweeney and
681 Thimann 1937; Thimann et al. 1992; Thimann and Biradivolu 1994; Waller and
682 Nick 1997; Wang and Nick 1998; Holweg et al. 2004). Treatment with BFA, a
683 fungal inhibitor of vesicle budding, caused, despite the presence of auxin, a rapid
684 bundling of microfilaments and shifted actin from the cytosolic fraction into the
685 microsomal fraction (Waller et al. 2002) depending on the dose of auxin and of
686 BFA. In parallel, BFA shifted the dose-response curve of auxin-dependent growth
687 to higher concentrations. In other words, BFA decreased auxin sensitivity *in sensu*
688 *strictu*, consistent with an actin-dependent transport of auxin-signalling components
689 such as auxin-efflux carriers. Again, a self-amplification loop emerges, consisting
690 of auxin-dependent organization of actin filaments and actin-dependent transport of
691 auxin-signalling components.

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A prediction from this model for the actin-auxin feedback loop would be that a bundling of actin should be followed by a reduction in the activity of polar auxin transport. This prediction is supported by the suppression of division synchrony in tobacco cell lines that overexpress mouse talin (Maisch and Nick 2007). However, to measure auxin transport directly, it would be necessary to administer radioactively labelled auxin to one pole of the cell file and to quantify the radioactivity recovered in the opposite pole of the file. This is not possible in a cell culture that has to be cultivated as suspension in a liquid medium. This approach would be feasible, however, in the classical graminean coleoptile system, where auxin transport can be easily measured by following the distribution of radioactively labelled IAA fed to the coleoptile apex. Transgenic rice lines were generated that expressed variable levels of the actin-binding protein talin (Nick P, Han MJ, An G, submitted). In those lines, as a consequence of talin overexpression, actin filaments were bundled to variable extent, and this bundling of actin filaments was accompanied by a corresponding reduction in the polar transport of auxin and gravitropic curvature (as a physiological marker that relies on the activity of auxin transport). When a normal configuration of actin was restored by addition of exogenous auxin, this restored auxin transport as well. This rescue was mediated by transportable auxin species, but not by 2,4-D, which lacks polar transport. With use of this approach, the causal relationship between actin configuration and polar auxin transport could be shown directly.

[Au7]

[Au8]

[Au9]



A further prediction of the actin–auxin feedback model is oscillations in transport activity, because auxin will, through the reorganization of actin, stimulate its own efflux such that the intracellular level of auxin will drop, which in turn will result in a bundled configuration of actin, such that auxin-efflux carriers will be sequestered, culminating in a reduced efflux such that auxin received from the adjacent cells will accumulate and trigger a new cycle. The frequency of these oscillations should depend on the dynamics of actin reorganization (around 20 min; Nick P, Han MJ, An G, submitted), and the speed of PIN cycling (in the range of 5–10 min; Paciorek et al. 2005) and is therefore expected to be in the range of 25–30 min. In fact, classic experiments on basipetal auxin transport in coleoptiles report such oscillations with a period of 25 min (Hertel and Flory 1968).

[Au10]

We therefore arrive at a model of a self-referring regulatory circuit between polar auxin transport and actin organization, where auxin promotes its own transport by shaping actin filaments. Thus, similar to the patterning of cell division, the actin-auxin oscillator seems also to be the pacemaker for the patterning of cell expansion.

7 Why Auxin - or Order Without a “Great Chairman” 728



Already in multicellular algae, a polar transport of auxin can be detected (Dibb-Fuller and Morris 1992; Cooke et al. 2002) and seems to play a role in the establishment of polarity (Basu et al. 2002), indicating that the central role of auxin in cell communication is evolutionarily quite ancient and had already been developed prior

733 to the colonization of terrestrial habitats. Why has evolution selected such a simple
734 molecule for such a central role in intercellular coordination?

735 Although we are far from providing a full answer to this question, it seems that the
736 answer is related to plant-specific features in the organization of signalling and devel-
737 opment: In animal development, cell differentiation is typically controlled by precise
738 and defined regulatory networks that are structured by predetermined hierarchies.
739 In contrast, plant cells are endowed with a pronounced developmental flexibility that
740 is maintained basically throughout the entire life span of a plant cell. Moreover, there
741 are hardly any fixed hierarchies – plant development does not know of such a thing
742 as a “Great Chairman” that assigns differential developmental pathways to the indi-
743 vidual cells. Plant cells rather “negotiate” their individual developmental fates in a
744 fairly “democratic” manner with hierarchies being created ad hoc by mutual interac-
745 tions. It seems that auxin is a central tool in these “negotiations”, because it repre-
746 sents a versatile tool to establish ad hoc hierarchies on the background of the high
747 degree of “cellular anarchy” and noise that is characteristic for plant development.

748 Why is plant development so “noisy”? The strong developmental noise seems to
749 be the tribute paid to indeterminate morphogenesis. The manifestation of the *Bauplan*
750 in an individual plant depends strongly on the environmental conditions encountered
751 during development. This developmental flexibility includes a rapid response of
752 cell expansion, complemented by a somewhat slower addition of morphogenetic
753 elements, such as cells, pluricellular structures and organs. This patterning process
754 can integrate signals from the environment, and must therefore be both highly
755 flexible and robust. More specifically, the patterning process has to cope with
756 signals that can vary over several orders of magnitude for the strength of the control
757 signal, and new elements have to be added such that the pattern formed by the
758 preexisting elements is perpetuated and/or complemented.

759 Plant sensing occurs in a rather diffuse manner – there are no such things as
760 eyes, ears or tongues; there are, instead, populations of relatively undifferentiated
761 cells that sense environmental cues and signals. Nevertheless, plant sensing is surpris-
762 ingly sensitive. When this high sensitivity of signalling is reached without special-
763 ized sensory organs, the individual cells must already be endowed with very
764 efficient mechanisms for signal amplification that are active already during the first
765 steps of the transduction chain. This strong signal amplification will inevitably
766 result in all-or-none outputs of individual cells. The efficient amplification of weak
767 stimuli on the one hand, with the simultaneous necessity to discriminate between
768 very strong stimuli of different amplitude, poses special challenges for plant signal-
769 ling. If all cells of a given organ were absolutely identical and homogeneous, even
770 an extremely weak stimulation would yield a maximal response of the whole
771 organ. It is clear that such a system would not have survived natural selection – the
772 amplitude of the output must vary according to variable amplitudes of the input
773 signal, because the plant has to respond appropriately to stimuli that vary in inten-
774 sity, even if these stimuli are strong. One way to reconcile the requirement for high
775 sensitivity with the requirement for a graded, variable output would be to assign the
776 two antagonistic tasks to different levels of organization: the high sensitivity to the
777 individual cells that perceive the signal; the graded, variable output to the population

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of cells (i.e. the tissue or organ) by an integration over the individual cell responses. 778
 But this works only when the sensory thresholds of individual cells differ over the 779
 population; in other words, when the individual cells are highly heterogeneous with 780
 respect to signal sensitivity and thresholds. 781

This heterogeneity was actually observed when photomorphogenesis was 782
 investigated on a cellular level for phytochrome-induced anthocyanin patterns in 783
 mustard cotyledons, a classic system of light-dependent plant patterning (Mohr 784
 1972; Nick et al. 1993) or for microtubule reorientation in coleoptiles triggered by 785
 blue light or auxin depletion (Nick 1992). Even adjacent cells exhibited almost 786
 qualitative differences although they had received the same dose of the signal. 787
 However, when the frequency of responsive cells in a given situation was scored 788
 and plotted against the strength of the stimulus, a highly ordered function emerged. 789
 Thus, the realm of individual cells was reigned over by chaos; order emerged only 790
 on the level of the whole organ. This highly stochastic, all-or-none type response of 791
 individual cells becomes especially manifest for an early response to a saturating 792
 stimulus or as the final result of weak induction (Nick et al. 1992). It thus appears 793
 that early signalling events are highly stochastic, when assayed at the level of 794
 [Au11] individual cells. These responses are not merely “noisy” because the flexible physiology 795
 of plant cells can tolerate this. These “noisy” responses rather represent an innate 796
 system property of plant signalling. However, this “noisy” inputs poses especial 797
 challenges for any ordering principle. 798

It might be these challenges that have rendered IAA a central integrator and 799
 synchronizer of plant development. A molecule that is easily transported through the 800
 acidic environment of the apoplast, but that is readily trapped in the cytoplasm and then 801
 has to be actively exported is ideally suited to convey lateral inhibition between 802
 neighbouring cells. It was sufficient to put the localization of the efflux transporter 803
 (whatever its molecular nature may be) under the control of auxin itself to reach a 804
 perfect reaction-diffusion system *in sensu* Turing (1952). On the intracellular level, 805
 this system is able to establish a clear cell polarity from even minute and noisy direc- 806
 tional cues. On the level of tissues, this cell polarity will generate patterns in a manner 807
 that meets the special constraints of plant development, i.e. noisy inputs as a conse- 808
 quence diffuse sensing and progressive addition of new elements to the pattern. Since the 809
 natural auxin IAA can enter the cell from any direction (because it can enter the cell 810
 even independently of import carriers such as AUX1), but will exit in a defined direc- 811
 tion defined by the localized activity of the efflux carrier, it can collect the input from 812
 several neighbours and focus this input into a clear directional output. It is this property 813
 that makes auxin a versatile and robust integrator for cell–cell communication. 814

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Author Queries

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Queries	Details Required	Author's Response
[AU1]	Please advise if "basis" is meant.	
[AU2]	Please advise if this should be "ARP".	
[AU3]	Kindly note that "Maisch J, Fischer L, Nick P, Submitted" is not found in the reference list. Please add this reference to the list or delete from the text.	
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