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REVIEW

# The endodermis—development and differentiation of the plant's inner skin

Julien Alassimone · Daniele Roppolo · Niko Geldner ·  
 Joop E. M. Vermeer

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**Abstract** Controlling external compound entrance is essential for plant survival. To set up an efficient and selective sorting of nutrients, free diffusion via the apoplast in vascular plants is blocked at the level of the endodermis. Although we have learned a lot about endodermal specification in the last years, information regarding its differentiation is still very limited. A differentiated endodermal cell can be defined by the presence of the “Casparian strip” (CS), a cell wall modification described first by Robert Caspary in 1865. While the anatomical description of CS in many vascular plants has been very detailed, we still lack molecular information about the establishment of the Casparian strips and their actual function in roots. The recent isolation of a novel protein family, the CASPs, that localizes precisely to a domain of the plasma membrane underneath the CS represents an excellent point of entry to explore CS function and formation. In addition, it has been shown that the endodermis contains transporters that are localized to either the central (stele-facing) or peripheral (soil-facing) plasma membranes. These features suggest that the endodermis functions as a polar plant epithelium.

**Keywords** Endodermis · Polarity · Casparian strip · Cell differentiation · Epithelium · *Arabidopsis thaliana*

## Abbreviations

AUX1	AUXIN RESISTANT1
AXR4	AUXIN RESISTANT 4
BOR1	BORON EFFLUX CARRIER 1
BOR4	BORON EFFLUX CARRIER 4
CS	Casparian strip
CSD	Casparian strip membrane domain
GFP	Green fluorescent protein
FM4-64	<i>N</i> -(3-triethylammoniumpropyl)-4-(6-(4-(diethylamino)phenyl)hexatrienyl)pyridinium dibromide
NIP5; 1	NOD26-LIKE INTRINSIC PROTEIN 5; 1
NPSN12	Novel plant snare 12
PEN3	Penetration resistant 3
PI	Propidium iodide
PIN	PINFORMED
PIS1	Polar auxin transport inhibitor sensitive 1
PM	Plasma membrane
PP2A	PROTEIN PHOSPHATASE 2A
SCR	SCARECROW
SHR	SHORT ROOT

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J. Alassimone · D. Roppolo · N. Geldner · J. E. M. Vermeer (✉)  
 Department of Plant Molecular Biology, University of Lausanne,  
 Quartier Sorge,  
 Lausanne 1015, Switzerland  
 e-mail: Joop.Vermeer@unil.ch

J. Alassimone  
 e-mail: Julien.Alassimone@unil.ch

D. Roppolo  
 e-mail: Daniele.Roppolo@unil.ch

N. Geldner  
 e-mail: Niko.Geldner@unil.ch

## Introduction

Every single cell in a higher plant is supplied with mineral compounds drawn from the soil by the roots and distributed to the aerial organs via the vascular tissues. Because of their immobility, terrestrial plants are fully dependent on their surrounding environment and must deal with the environmental conditions they live with. The roots have the

fundamental role of being the interface involved in nutrients intake. Yet, they must at the same time provide an efficient boundary against external biotic stresses, such as pathogenic microorganisms or abiotic stresses, such as excessive ion concentrations.

External compounds can reach the stele and be distributed throughout the plant following two routes; they either move from one cell to the other via the symplastic pathway or progress between cells via the apoplast (Marschner 1995) (Fig. 1).

In the symplastic pathway, the plasma membrane (PM) acts as the first soil/plant interface by selecting ion uptake through specific transporters (Marschner 1995). Passive diffusion through the membrane or unspecific uptake may lead to the unintended entrance of compounds, but transporters regulating active export are known to be present in the epidermis and can counteract the presence of undesired molecules (Miwa et al. 2007). Controlling external compound entrance via the apoplastic space is more complicated since free diffusion occurs within cell walls. Thus, in order to allow an efficient and selective sorting of nutrients, an apoplastic diffusion barrier is present in the endodermis of all vascular plants, making passage through endodermal cells a mandatory step for molecules to reach the stele.

## Endodermis specification

Endodermis specification has been intensively studied over the last years. Most of our knowledge regarding this process comes from research carried out in the model

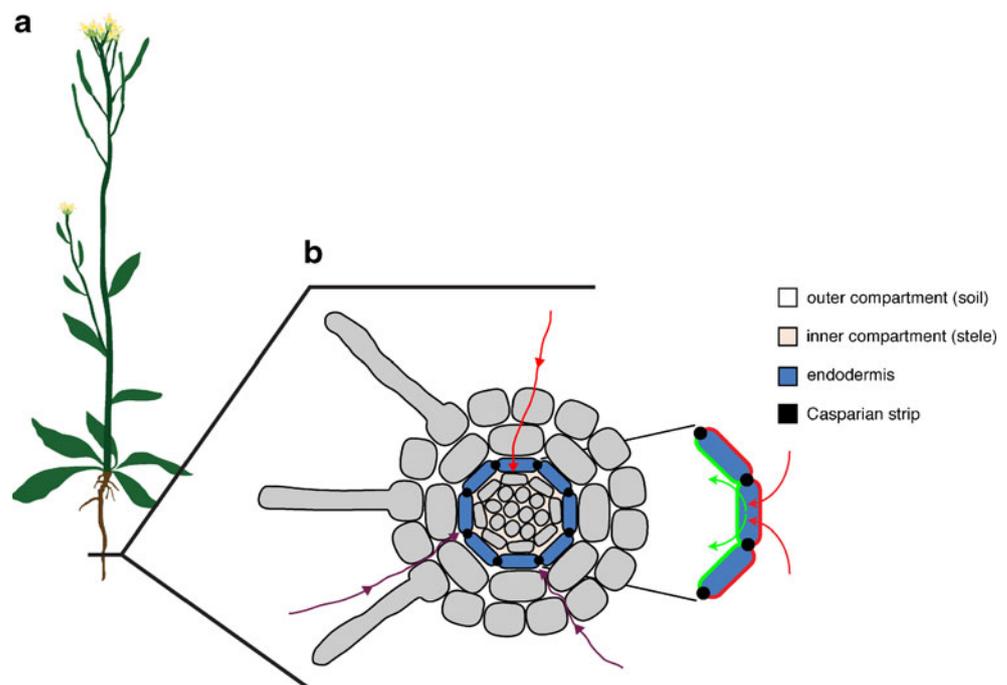
plant *Arabidopsis thaliana*. In roots, cell types and developmental stages are easily recognizable: tissues are organized in concentric rings along the radial axis while developmental zones can be distinguished along the longitudinal axis. The cell specification of initials leading to endodermal cells is regulated by the interaction between two transcription factors, SHORTROOT (SHR) and SCARECROW (SCR), that together trigger the periclinal division of cortex–endodermal initials leading to the formation of the endodermis and the cortex (Helariutta et al. 2000). This process requires the movement of SHR from the stele, where it is expressed, into the endodermis; here, SHR interacts with SCR leading to the transcription of target genes and eventually determination of endodermal cell identity (Benfey and Scheres 2000; Cui et al. 2007; Di Laurenzio et al. 1996; Gallagher et al. 2004; Helariutta et al. 2000; Nakajima et al. 2001; Sozzani et al. 2010; Wysocka-Diller et al. 2000).

Although specification of endodermal initials is a well-understood process, we are still completely in the dark about the molecular events that determine endodermis differentiation. This review will focus on cell differentiation by addressing the development, early differentiation, and CS formation of endodermal cells.

## The endodermis: not just another cell layer

It has been a long time since the first description and suggestion of a diffusion barrier present in the “plant inner skin” by Robert Caspary, which is nowadays known as the

**Fig. 1** The endodermis, and not the epidermis, is the barrier for the extracellular diffusion of molecules. **a** Schematic drawing of an *Arabidopsis* plant and **b** a detailed view of a cross-section of the root. Purple lines show the apoplastic route molecules can take until they are blocked by the CS. The inset shows an enlargement of a few endodermal cells with proposed central (green) and peripheral (red) localized transporters that facilitate the loading of nutrients in the inner space. Red arrows indicate symplastic route of uptake of compounds



Casparian strip (CS) (Caspar 1865). This structure refers to a modification that occurs by impregnation with ligno-suberic material of the anticlinal primary cell wall of endodermal cells, resulting in the formation of an equatorial belt-like structure (Fig. 1). The CS occupancy is normally about one third of anticlinal walls but can increase in size when exposed to environmental stresses, suggesting a certain plasticity of this structure (Karahara et al. 2004). The CS is the primary developmental stage of endodermal differentiation into a “border sheath”, and it is generally followed by a suberin deposition, forming suberin lamellae that eventually coat the entire cell (for a review, see Enstone et al. 2003). The coated cell is still connected to the surrounding cells by plasmodesmata (Clarkson et al. 1987), although plasmodesmata are not observed in the region of the CS itself (Bonnett 1968). In some plant species, U-shaped tertiary cell walls have been described in the very late developmental stage of endodermal cells (for a review, see Enstone et al. 2003). Other examples of apoplastic barriers featuring CS-like structures exist in the root exodermis, in the needles of Pinaceae, and in specific cells involved in nectar secretion (Enstone et al. 2003; Liesche et al. 2011).

The CS is an endodermal differentiation feature considered to be crucial for plant development and survival. Undeniably, it provides a diffusion barrier between the apoplastic space facing the soil and the innermost apoplastic space, in the stele. The efficiency of the CS to act as a diffusion barrier has been highlighted by several studies demonstrating that the transport of dyes (Moon et al. 1986; Rufz de Lavison 1910; Weerdenburg and Peterson 1984) or ions (Peterson 1987; Robards and Robb 1974; Rufz de Lavison 1910; Singh and Jacobson 1977) but also heavy metals (Nagahashi et al. 1974; Robards and Robb 1972) are blocked and remain in the apoplast of cortical tissues. As a direct consequence of the establishment of this barrier, a directional and selective transport of nutrients should occur across the endodermis (Fig. 1b). Indeed, influx and efflux carriers have been localized to the central (stele-facing) and peripheral (soil-facing) sides of the endodermal plasma membrane (Alassimone et al. 2010; Ma et al. 2006, 2007; Takano et al. 2010). Thus, the CS isolates the stele's apoplastic space and distinct domains are organized in the endodermal plasma membrane. Beyond selective nutrient uptake, this is beneficial to prevent extensive ion backflow from the stele to the soil, and to maintain root pressure (Luttge and Laties 1966; Peterson et al. 1993). The latter is known to be important in sieve transport under low transpiration and in preventing xylem embolism in case of xylem cavitations (Tyree and Sperry 1989). Currently, the role of the endodermis in defense against biotic stresses remains unclear despite evidence indicating that the CS is resistant to cell-wall-degrading enzymes that are generally

used by pathogens to degrade primary cell walls (Schreiber et al. 1999). An accumulation of pathogens at this apoplastic border has not been reported so far. Interestingly, juvenile root knot nematodes avoid the CS on their way to the vasculature: after penetration of the root epidermis, they move through the cortical apoplast towards the root tip and finally penetrate the vascular bundle at the meristematic or elongation zone, where CS are not yet established (Williamson and Gleason 2003).

### Endodermis differentiation

In the last decades, several reports have studied CS structure and function of already differentiated endodermal cells, but only recently the developmental sequence of events that leads to this differentiation has been described in *Arabidopsis thaliana* (Alassimone et al. 2010). In this study, molecular and histochemical methods were used to highlight several differentiation events: propidium iodide was used as an apoplastic tracer and revealed the establishment of a functional apoplastic barrier; CS impregnations were directly observed by visualizing their auto-fluorescence; and physical attachment of the PM to cell wall was observed after plasmolysis, while diffusion in the PM was evaluated by using the fluorescent lipophilic styryl dye FM4-64. The combination of these techniques allowed the identification of a membrane domain where free diffusion from one side to the other of the PM was prevented. This PM domain is formed underneath the CS and is referred to as the “Casparian strip membrane domain (CSD)”. After plasmolysis, the CSD remained attached to the cell wall, which is in agreement with the hypothesis that a protein-like structure may scaffold the CSD to the CS (Alassimone et al. 2010; Behrisch 1926; Bonnett 1968; Karahara and Shibaoka 1992).

As suggested, cell wall impregnations and PM attachment might be driven by protein scaffolding at the exact position of the CS, as previously suggested by ultrastructural studies (Bonnett 1968). Moreover, this protein scaffold may represent the barrier that prevents free diffusion of FM4-64 and plasma membrane proteins that are also excluded from the CSD (Alassimone et al. 2010). The establishment of a protein exclusion zone, the appearance of membrane–cell wall attachment sites, and the deposition of the CS all occur in a narrow time frame, while PI diffusion is blocked slightly later, suggesting that formation of a functional apoplastic barrier is a late differentiation step.

Therefore, several events can be observed during the development of mature endodermal cells: some of these events may be linked and occur concomitantly, while other seems to follow a step-by-step process (Alassimone et al. 2010).

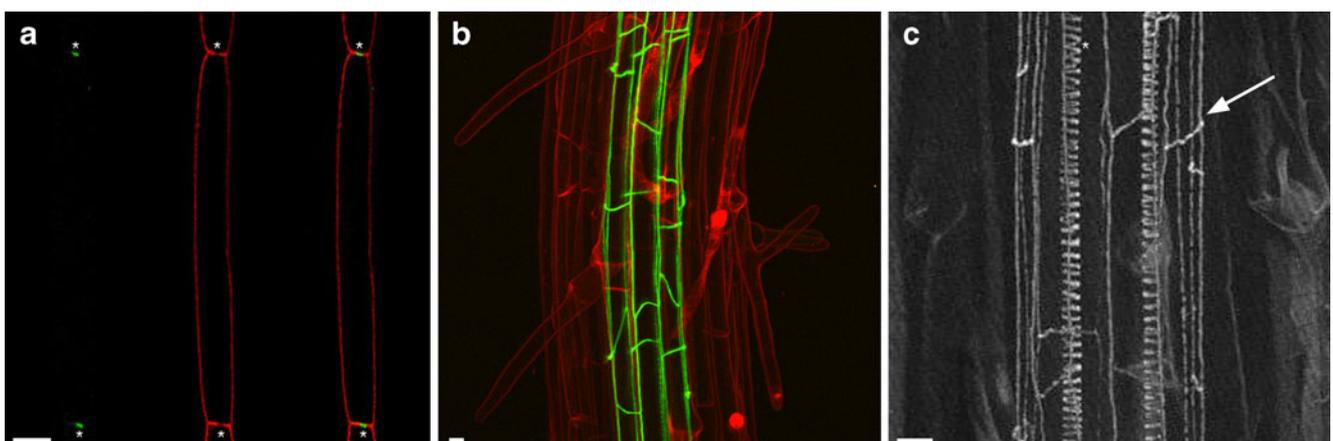
## A novel protein family localizes to the Casparian strip domain

Recently, it has been reported that the plasma membrane at the CS is inaccessible to plasma membrane proteins, which allowed defining this “Casparian strip membrane domain” (CSD) as a zone of protein exclusion. Although this important feature has never been described before, another group recently reported that the auxin efflux carrier PIN3 is also excluded from the CSD (Alassimone et al. 2010; Ding et al. 2011).

It can be hypothesized that the spatial separation of the two lateral membrane domains is mediated by a tight protein scaffold that would also cause the general exclusion of membrane proteins from the CSD. This hypothesis is supported by ultrastructural analysis of the CS (Bonnett 1968). Until recently, we were completely in the dark regarding the proteins that might localize to this exclusion zone. However, a recent publication describes the identification of a novel protein family consisting of five members that localize precisely to this exclusion zone (Roppolo et al. 2011). These proteins were named CASP1-5 (Casparian Strip membrane domain Protein). CASPs are 4-transmembrane spanning proteins that initially localize to the plasma membrane, but prior to the development of the CS localize to the exclusion zone (Fig. 2a). CASP transcription starts ~2 cells prior to the CSD formation, which matches the time it takes from its initial protein accumulation and final localization. This localization appears about five cells prior to the establishment of a functional diffusion barrier. This means that the CASPs are early markers for CSD and CS formation. In other words, the CASPs mark the differentiation of endodermal cells into

CS-containing endodermal cells (Roppolo et al. 2011). Their localization to the CSD domain prior to the establishment of the CSD implies an early role for the CASP proteins in CSD formation. Moreover, Roppolo et al. (2011) observed that in a *casp1;casp3* double mutant, the CS appeared to be disorganized and more fluorescent compared to the CS in wild-type plants, although it still functions as a barrier. This observation does suggest a role for the CASP proteins in regulating cell-wall modification needed for the formation of the CS. It also shows that the other remaining CASP proteins can sufficiently compensate for the loss of CASP1 and CASP3 to maintain the diffusion barrier. It is likely that higher order *casp* mutants are needed to reveal the exact role of CASPs in CS formation.

When tagged with the green fluorescent protein (GFP), CASP proteins show a similar net-like localization as the autofluorescence of the CS reveals (Fig. 2b, c). In addition, it has been shown that once the CASP proteins get localized into the CSD, they become completely immobile as compared to their initial plasma membrane localization (Roppolo et al. 2011). Immuno-electron microscopy experiments on plants expressing CASP1-GFP under the control of its own promoter showed that CASP1 localizes to a spatially restricted domain in the plasma membrane that is tightly aligned to the CASP1 domain in the adjacent plasma membrane of the next cell, separated by the CS cell wall. It is of interest to note that the CASP proteins seem only able to form the CSD domain in endodermal cells as ectopic expression results in only plasma membrane localization or accumulation in aggregates. This suggests that there is a specific signal in the endodermis that triggers the accumulation of CASP proteins in the CSD. How CASP proteins change their localization from an evenly



**Fig. 2** CASP proteins localize to the CS domain and mediate CS formation in *Arabidopsis*. **a** Confocal image showing a medial longitudinal section of a differentiated endodermal cell co-expressing CASP1-mCherry (green) and YFP-NPSN12 (red). Note that CASP1 localizes to the middle of the apical and basal membrane, precisely labeling the depletion zone, highlighted by the asterisks. **b** Maximal

projection of an image stack taken from a plant expressing CASP1-GFP (green). Cell outlines are stained with PI (red). CASP1-GFP forms a net-like structure encompassing the endodermis. **c** Maximal image projection of a cleared *Arabidopsis* root, showing the CS (arrow) and xylem autofluorescence (asterisk). Bars=10  $\mu$ m

distributed to a highly localized one at the CSD and how they align themselves between cells is currently not known.

Initially, CASP proteins are sensitive to treatment with the fungal toxin Brefeldin A (BFA) that affects endocytic recycling, but once localized to the CSD they are no longer sensitive to BFA and they also do not seem to turn over anymore (Roppolo et al. 2011). In addition, it was shown that CASP proteins can interact with each other, and this might result in the formation of a large, tightly organized protein complex capable of preventing diffusion of membrane proteins through this domain and providing the scaffold for the formation of the CS. The identification of the CASP proteins and the use of the fluorescent CASP protein fusions now open up a new field of research and will allow us to get more fundamental insights into how the CS is made and what roles it plays during plant development. In addition, now that the first proteins localizing to the CSD have been identified, it will facilitate the identification and functional analysis of other proteins that make up the CSD. We hope that the identification of the CASPs will cause a similar boost in our understanding of CS formation as the identification of the first tight junction-localized protein ZONULA OCCLUDENS-1 (ZO-1) did for research in animal epithelia.

Since the CSD defines two distinct plasma membrane domains, proteins confined to one lateral domain should not be able to diffuse into the opposite domain. Under these conditions, several influx or efflux carriers might be localized either on the central (stele-facing) or peripheral (soil-facing) domains of endodermal cells. Hence, studying the endodermis should allow the investigation of this polarity in a relevant context.

### Plasma membrane organization in plant roots: existence of distinct domains

Research on plant polarity has been intensively focused on the auxin-transport-related proteins, the PINFORMED (PIN) or AUXIN RESISTANT 1 (AUX1) proteins, that display an apical–basal (AB) polarity (for review, see Feraru and Friml 2008; Grunewald and Friml 2010).

In addition to AB polarity, some proteins have been shown to exhibit central–peripheral (CP) polarity as shown for silicon efflux and influx carriers in rice (Ma et al. 2006, 2007). In *Arabidopsis*, the BORON EFFLUX CARRIER 4 (BOR4) and the boron influx carrier NOD26-LIKE INTRINSIC PROTEIN 5;1 (NIP5;1) localize to the peripheral PM domain (Miwa et al. 2007; Takano et al. 2006). In contrast, the BORON EFFLUX CARRIER 1 (BOR1) is observed at the central PM domain (Alassimone et al. 2010; Takano et al. 2010). The plant pathogen-defense-related and hormone precursor transporters PEN3/PDR8/ABCG36 and

PIS1/PDR9/ABCG37 are also exclusively localized to the peripheral PM domain (Langowski et al. 2010; Strader and Bartel 2009). So far, it has been shown that BOR1, NIP5;1, and PIS1 display polar localization in endodermal cells, although PIS1 was ectopically expressed (Alassimone et al. 2010; Langowski et al. 2010; Takano et al. 2010) (Fig. 3b). It has also been shown that these markers, when ectopically expressed, exhibited polar localization in epidermal and cortical cells (Alassimone et al. 2010; Langowski et al. 2010; Takano et al. 2010).

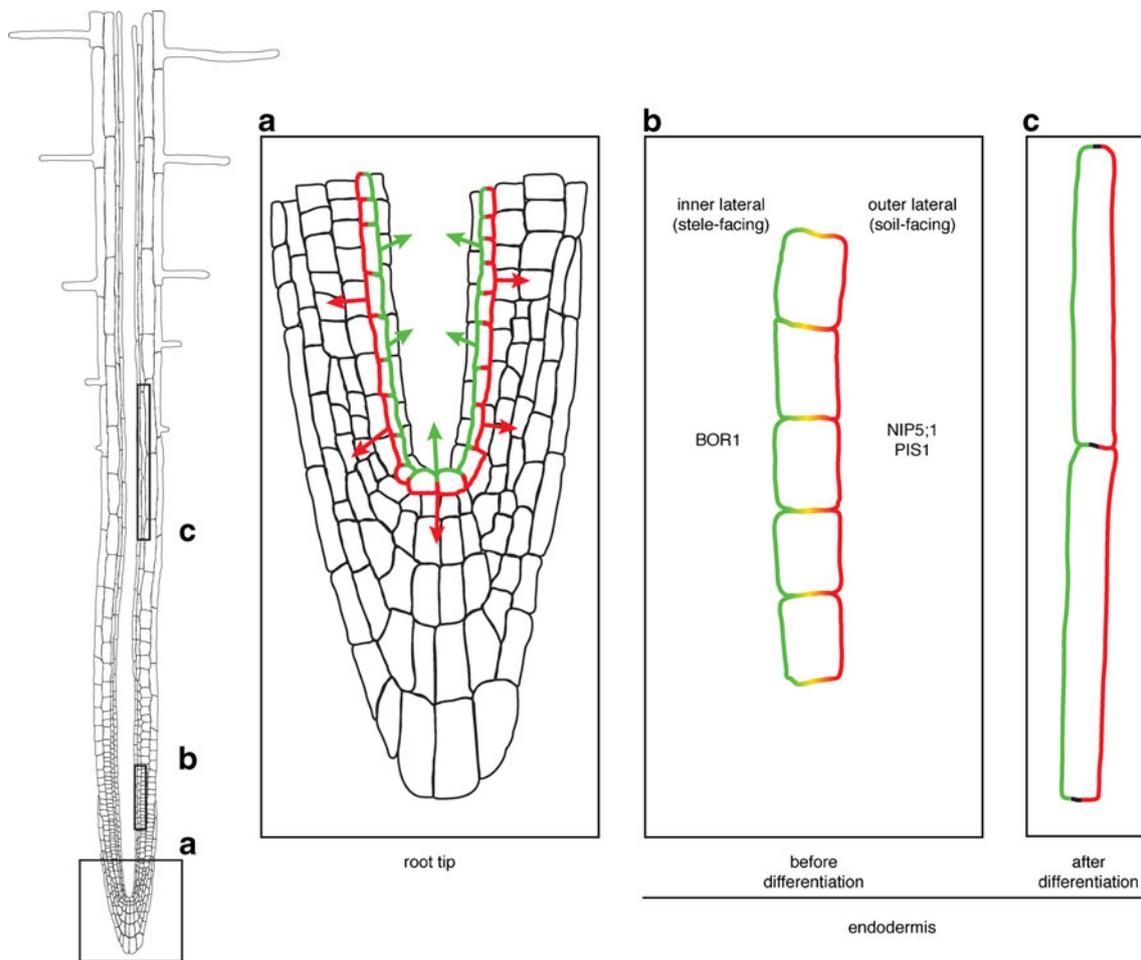
Our hypothesis regarding CP-polarity function is that it is needed to direct nutrient flux from the soil to the stele, but when nutrients reach the stele, CP polarity is no longer required. The sequestration of nutrients in the vascular cylinder, by the virtue of the CS, might make their directional delivery obsolete once inside the stele. The fact that also non-differentiated cells exhibit CP polarity also fits with this hypothesis. In fact, nutrients may well not solely move through symplastic or apoplastic routes but can also use a combination of both (Fig. 4). Interestingly, our hypothesis is supported by the fact that ectopically expressed PIS1 seems to lose its CP polarity inside the stele (Langowski et al. 2010).

As the CP polarity seems to be organized with respect to the stele position (Alassimone et al. 2010; Langowski et al. 2010; Takano et al. 2010) (Fig. 3a), it is tempting to speculate that a mobile signal from the stele might regulate CP polarity.

### Do AB polarity and CP polarity share common mechanisms?

Intuitively, two mechanisms could explain polar localization; either proteins are actively targeted to a defined domain or they are randomly secreted and specifically removed from the opposite domain (for a review, see Geldner 2009). In the absence of lateral diffusion barriers, continuous endocytosis should be required for the maintenance of polarity, even if non-localized and constitutive, simply to counteract the inevitable dissipation of polarity by lateral diffusion of proteins (Valdez-Taubas and Pelham 2003). Up to now, the major insights underlying establishment and maintenance of AB polarity in plant roots have been based on research studying the PIN family of auxin-efflux carriers (Feraru and Friml 2008; Grunewald and Friml 2010).

It is known that the basal localization of PIN1 is dependent on endocytic recycling mediated by the ADP-ribosylation factor guanine nucleotide exchange factor (ARF-GEF) GNOM as PIN1 polarity is strongly affected in *gnom* mutants or by inhibiting GNOM activity by the BFA (Geldner et al. 2003). In addition, it has been shown



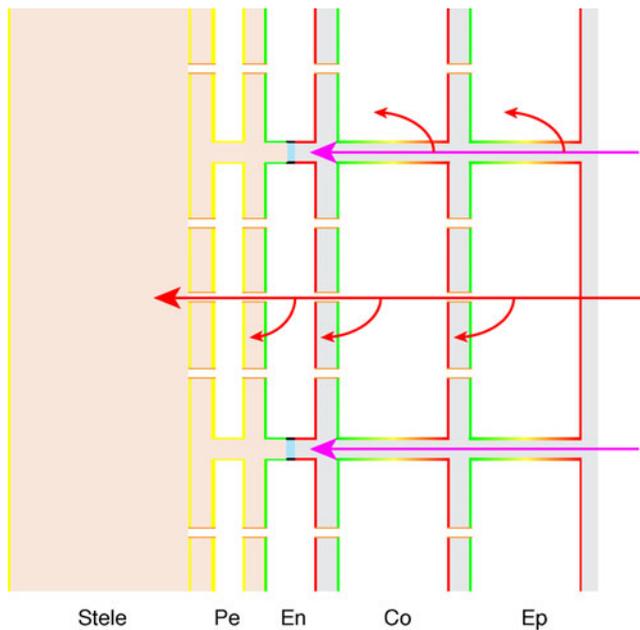
**Fig. 3** Polarity establishment in the endodermis precedes CS development. Schematic drawing of an *Arabidopsis* root. Boxed regions are enlarged in (a)–(c) to highlight endodermal polarity establishment. **a** Root tip of *Arabidopsis* displaying already stele-facing (green) and soil-

facing (red) polarity. **b** Zoom-in of undifferentiated endodermal cells showing an intermixing of the two polar lateral domains (yellow). **c** Zoom-in of differentiated endodermal cells. The establishment of CSD (black) results in a strictly separated central and peripheral polarity

that PIN1 is initially asymmetrically secreted and that subsequent polarization involves AtRabF2b/ARA7-mediated endocytosis and/or recycling. However, apical localized proteins like PIN2 and AUX1 seemed to be hardly affected, suggesting that there is at least another mechanism that regulates apical polarity (Dhonukshe et al. 2008). Over the years, several additional mechanisms have been shown to act in the internalization and cycling of PIN proteins (Furutani et al. 2011; Kitakura et al. 2011; Naramoto et al. 2010). For example, sterol composition of the PM, the cytoskeleton, and cell-wall anchoring have been suggested to be important for PIN polarity (Feraru et al. 2011; Geldner et al. 2001; Heisler et al. 2010; Kleine-Vehn et al. 2008; Men et al. 2008).

Although only few reports have currently addressed this question, it seems that the mechanism underlying CP-polarity establishment and maintenance differs from that underlying AB polarity. It has been shown by Fluorescence Recovery After Photobleaching (FRAP) experiments that

PIS1 seems to exhibit a polar exocytosis to the peripheral plasma membrane domain, whereas FRAP studies on PIN1 revealed non-polar exocytosis (Dhonukshe et al. 2008; Langowski et al. 2010). Treatments with BFA had no effect on the polar localization of BOR1, NIP5;1, or PIS1. However, BFA treatment did result in the re-localization of these proteins into endosomal aggregates, indicating that they are partially trafficked through BFA-sensitive endosomes (Alassimone et al. 2010; Langowski et al. 2010; Takano et al. 2010). In addition, it was shown that PIS1 was still correctly localized in the *gnom* mutant (Langowski et al. 2010). This does not mean that endocytic recycling is not as important for CP polarity as is the case for AB polarity, but just that GNOM seems not to be crucial for CP polarity establishment and maintenance. Additional studies are required to identify the molecular mechanisms mediating CP polarity. One intriguing observation pointing to an important role for membrane trafficking in CP polarity is that treatment with the PI3-kinase inhibitor wortmannin



**Fig. 4** Different routes for nutrient uptake by plant roots. Schematic representation of an *Arabidopsis* root section showing different routes of compound uptake. From the epidermis to the endodermis there are stele- (green) and soil-faced (red) carriers which could provide a flux of molecules to the stele. Once passed the endodermis, this intrinsic lateral polarity seems lost (highlighted by a yellow-colored plasma membrane). Red arrows indicate the symplastic route and the purple arrows the apoplastic route. Of course, transport can also consist of a combination of the two pathways. Plasmodesmata are shown in orange, CSD in black, and the CS in blue. For clarity, plasmodesmata present in the anticlinal walls have been omitted. *Pe* pericycle, *En* endodermis, *Co* cortex, *Ep* epidermis

(WM) induces different effects on NIP5;1 compared to BOR1 in endodermal cells. High WM concentration does not affect BOR1 polarization, but does depolarize NIP5;1. Interestingly, this only occurred in endodermal cells where the CS was not yet established (Alassimone et al. 2010). This depolarization is thought to be a consequence of a block of PM internalization by WM, allowing lateral diffusion to distribute NIP5;1 to the whole PM domain in the absence of the CSD. This also suggests a role for the CS in lateral polarity maintenance by providing a PM diffusion barrier and resulting in the confinement of the proteins to one lateral PM domain.

In addition to endocytic recycling, it is known that phosphorylation/dephosphorylation processes involving the Ser/Thr protein kinase PINOID (PID) and the protein phosphatase 2A (PP2A) act in PIN polarity establishment (Friml et al. 2004; Michniewicz et al. 2007). However, PID and PP2A do not seem to be required for peripheral polarity (Langowski et al. 2010). Besides this, it has been shown that the ER-localized AUXIN RESISTANT 4 (AXR4), needed for the correct polar localization of AUX1, is also not required for BOR4, PIS1, or PEN3 polarity (Langowski

et al. 2010) and that the sterol composition of the plasma membrane has no effect on CP polarity (Langowski et al. 2010).

No obvious correlation between lateral polarity and the cytoskeleton has been reported so far. Chemical treatments or imaging of the cytoskeleton using GFP probes have not revealed any link to polarity establishment (Alassimone et al. 2010; Langowski et al. 2010; Takano et al. 2010). Disrupting the actin polymerization revealed that PIS1 and PEN3 have a tendency to accumulate in intracellular aggregations (Langowski et al. 2010). Very similar intercellular aggregations of a non-polar PM marker protein could also be observed after chemically induced actin depolymerization in endodermal cells (Alassimone et al. 2010). Thus, intracellular aggregations observed after actin depolymerization might not be specific to targeting of polar proteins, but rather a result of a general interference with membrane trafficking.

Although only a few proteins exhibiting CP polarity have been identified, there is already evidence indicating differences between some of them. For example, independent of polarity-related recycling, BOR1 endocytosis and degradation is regulated by boron availability whereas NIP5;1 and BOR4 are not (Miwa et al. 2007; Takano et al. 2005, 2010). Although both are partially BFA sensitive, BOR1 seems to be more sensitive to BFA than NIP5;1 (Takano et al. 2010). Ubiquitination has been reported to be required for BOR1 sorting but seems not to be required for polar localization (Kasai et al. 2011). The lysine residue shown to be responsible for the ubiquitination-dependent vacuolar sorting in BOR1 seems to be absent from BOR4 (Kasai et al. 2011). Moreover, tyrosine motifs have been shown to be necessary for the proper polar localization of BOR1 (Takano et al. 2010). It seems unlikely that tyrosine motifs are important for NIP5;1 localization since the two potential tyrosine-based signals are predicted in transmembrane and extracellular regions (Takano et al. 2010). It is clear that we need to isolate additional proteins displaying CP polarity to be able to start addressing the mechanistic framework required for this cellular process.

#### A function for the CSD in endodermal polarity?

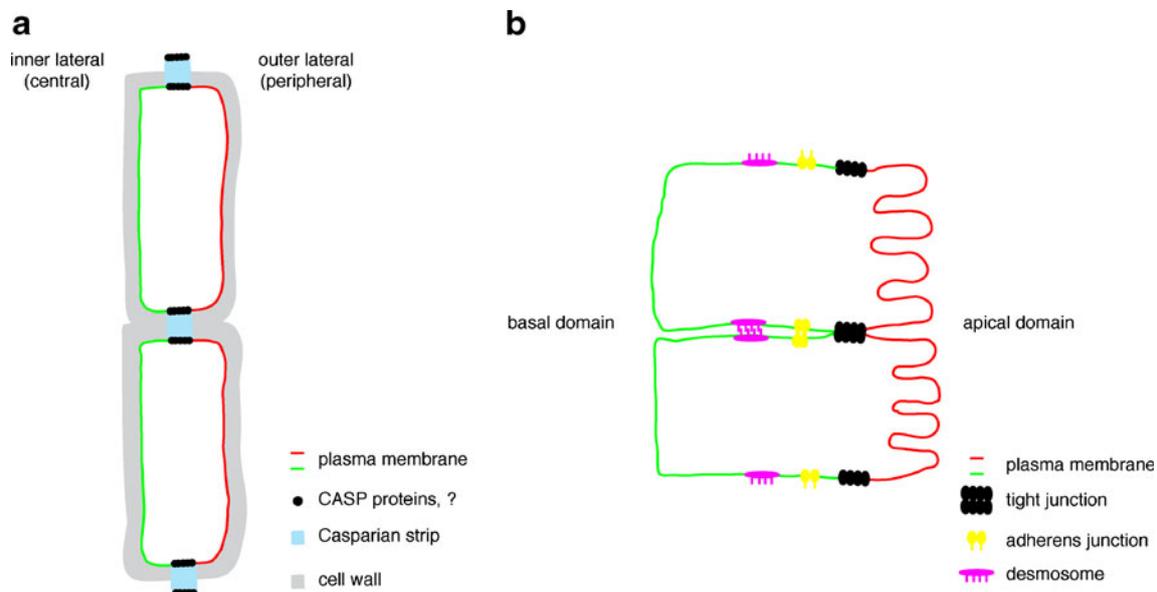
Endodermal cells are able to organize lateral polarity even before the partitioning of their PM into two lateral domains (Alassimone et al. 2010). Furthermore, lateral polarity is observed in cells lacking the CSD such as epidermal or cortical cells (Alassimone et al. 2010; Langowski et al. 2010; Takano et al. 2010). The presence of a functional barrier seems not to be required for polarity establishment or maintenance, but is effective for a physical separation of the two lateral membrane domains. Indeed, polar proteins

in differentiated endodermal cells are no longer intermixed, but confined to one lateral side of the cell (Alassimone et al. 2010) (Fig. 3c). This suggests that the observed localization of polar proteins in the anticlinal PM of undifferentiated endodermal cells could be due to lateral diffusion. In addition, Langowski et al. proposed a dose-dependent effect on polarity as the localization of PIS1 became less pronounced when overexpressed (Langowski et al. 2010), suggesting that without the physical restriction of lateral domains by the CS, a possible saturation of the machinery involved in polarity maintenance could occur resulting in inefficient confinement of polar proteins.

### The endodermis as a polar epithelium of plants

The endodermis provides a barrier that separates the inner (stele) tissue from the outer (soil) interface and thus contains stele-facing and soil-facing plasma membrane domains that are separated by the CSD, to regulate uptake of nutrients (Figs. 3 and 4). In addition, the barrier made up out of the CSD and CS bears functional similarity with mammalian tight junctions as it separates two membrane domains and forms a barrier that seals off the outer from the inner tissues. Although plants lack homologues of proteins known to be required for the establishment of polarized epithelia in vertebrates, such as PAR proteins, CLAUDINS, OCCLUDINS, ZO proteins, and others (Cereijido et al.

2004; Martin-Belmonte and Mostov 2008; Tsukita et al. 2009), the endodermis does have all the functional features of a polar epithelium. While the plant epithelium has only one barrier complex, the CS, the mammalian epithelial junctional complex consists of tight junctions, adherens junctions, and desmosomes. Since the tight junction is the most apical complex and it forms a tight seal between adjacent cells thereby separating the external and internal environment, it bears most similarity to the CSD and CS (Gupta and Ryan 2010; Steed et al. 2010; Bazzoni and Dejana 2004; Zeeb et al. 2010; Forster 2008). Indeed, although the CASP proteins do not show any obvious sequence similarity when compared with CLAUDINS, they do share similarity in overall structure (both 4-TM proteins) and size (Grebe 2011). Moreover, it has been reported that similar to CASPs, the CLAUDINS show very limited diffusion in tight junctions, whereas other tight junction localized proteins such as OCCLUDINS and ZO-1 show much more mobility (Shen et al. 2008). CLAUDINS are viewed as the core constituents of tight junctions, a function that is also suggested for CASPs (Roppolo et al. 2011; Shen et al. 2008). Figure 5 shows a schematic representation of the endodermal and epithelial cells. The most obvious difference between the CS and tight junctions is the fact that in the CS there seems to be a “no-touch policy”; the two adjacent CSD are separated by the cell-wall modification known as the CS. Hence, the CASPs of one endodermal cell will never be in contact with the CASPs



**Fig. 5** The endodermis as a polar epithelium of plants. The endodermis bears some overall similarity with polar epithelium of vertebrates. The Casparian strip forms a barrier that will separate the stele from the soil interface. The CASP proteins are suggested to be important for the formation of this barrier. In epithelia, the junctional complex is made up by tight junctions, adherens junctions, and

desmosomes. The tight junctions form the barrier that seals of the paracellular space from the lumen and the inside tissue, resembling the function of the Casparian strip. The big difference is that tight junctions are mediated by protein–protein contacts, whereas the CS is a cell-wall modification that does not depend on protein–protein interactions between the two adjacent cells

of the adjacent endodermal cell. This is in strong contrast to tight junctions, whose function is based on direct protein–protein interactions of tight junction complex-localized proteins between two adjacent cells (e.g., CLAUDINS) (Bazzoni and Dejana 2004; Forster 2008; Gupta and Ryan 2010; Shen et al. 2008; Steed et al. 2010; Zeeb et al. 2010). Tight junctions are viewed as dynamic structures that function as selective barriers allowing certain molecules to pass through the paracellular space whereas others are blocked (Steed et al. 2010). At the moment, we do not know whether the CS can function in a similar way, i.e., as a kind of molecular sieve that allows selective passage of small molecules (e.g., water). Future research will be able to provide some answers to this. However, even if there is a functionally relevant selectivity of the CS toward certain small molecules, it is very probable that this selectivity is not subject to dynamic regulation as can be seen for the permeability and selectivity of animal tight junctions. This is because the more complex structure of the CS is not solely based on protein interactions but represents a localized modification/impregnation of the plant cell wall with hydrophobic polymers that probably cannot easily be altered in response to environmental stimuli.

### Future prospects

Although described almost 150 years ago by Robert Caspary, until recently we have not gained much knowledge underlying CS formation in plants and in which biological process(es) it is involved. Now, with the discovery of the CASP proteins, we finally can start to unravel how this barrier gets established. For example, genetic screens on CASP1::CASP1–GFP plants might reveal how CASPs are regulated, but might also reveal how they get localized so precisely to the CSD. In parallel, expression-based analysis, as successfully used for the CASP proteins, together with proteomics could be used to identify additional proteins that localize to the CSD and CS.

The generation of higher order CASP mutants, together with newly identified mutants that are required for barrier formation, will be instrumental to finally obtain genetic evidence for the biological roles of the CS. What will be the consequences of alterations of CS on drought, nutrient stress resistance, or the defense against soil borne pathogens?

Another interesting question is whether plants also are capable of locally modifying or altering their CS during lateral root formation. In order to emerge, lateral roots first need to pass the fully differentiated and CS-containing endodermis. Although our knowledge about lateral root formation has greatly advanced, surprisingly little is known about what happens to the endodermis during this process.

Some older reports on studies on other species speculate that this seems to be a regulated process (Bell and McCully 1970; Bonnett 1969; Karas and McCully 1973). Therefore, it might be that plants have devised a way to locally modify their CS to facilitate lateral root emergence. Besides deciphering the biological roles of the CS, the endodermis now also provides a beautiful system to study cell differentiation with powerful genetic and cell biological tools. We are still far away from a mechanistic understanding of CS formation, but the discovery of the CASP proteins and CS mutants provides us with the perfect tools to start investigating these processes.

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