

The Role of Auxin in Abscission of Organs and Tissues

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Cover: Abscission zone of a *Populus* leaf with GUS staining
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Abstract

Most deciduous trees drop their leaves before winter, a process which is referred to as leaf abscission. Leaf abscission is thought to be regulated by the action of auxin and ethylene. In order to test the function of auxin in leaf abscission, an experimental system in *Populus* was established to induce leaf shedding synchronously under controlled greenhouse conditions. Exogenous auxin and an auxin transport inhibitor delayed the abscission of dark-induced leaves and a new auxin response maximum preceded the formation of an abscission zone. The analysis of microarray results revealed that several genes encoding auxin transporters were strongly down-regulated during abscission, suggesting their involvement in the formation of the auxin maximum in the leaf axil. In ethylene-insensitive trees, leaf abscission could be delayed by the application of auxin and ethylene signaling was not required for the regulation of gene expression of auxin transporters during abscission. Thus, auxin and ethylene act partly independently of each other on leaf abscission in *Populus*.

In order to study the effects of auxin on cell separation, isolated from its action on the development of an abscission zone, we examined root cap abscission in *Arabidopsis*. An auxin response gradient, spanning the root cap, was found to be established prior to the separation of the outermost root cap layer. Inhibition of polar auxin transport abolished the auxin response gradient in the root cap and disrupted abscission. Intriguingly, auxin efflux carriers of the PIN family were not expressed in the cell layer proximal to the abscising layer indicating that the outermost columella tier is disconnected from the auxin source in the quiescent center.

A *Populus* homolog of the *Arabidopsis* WALLS ARE THIN1 (WAT1) was among the most strongly regulated genes during abscission. We found that WAT1 localizes to the tonoplast and facilitates auxin export from the vacuole. Whereas, WAT1-mediated auxin homeostasis is needed for secondary wall deposition, *wat1* mutants do not display any phenotype related to abscission.

While auxin gradients have been implicated in various growth-related processes our work provides novel data in support of a regulatory role of distinct auxin maxima and minima in organ and tissue abscission.

Keywords: *Populus*, auxin, ethylene, abscission, cell separation, PIN proteins, *Arabidopsis*, root cap, auxin transport.

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It's fine to celebrate success but it is more important to heed the lessons of failure.

- Bill Gates

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List of Publications

This thesis is based on the work contained in the following papers, referred to by Roman numerals in the text:

- I **Xu Jin**, Jorma Zimmermann, Andrea Polle and Urs Fischer (2015). Auxin is a long-range signal that acts independently of ethylene signaling on leaf abscission in *Populus*. *submitted*.
- II **Xu Jin**, Urs Fischer. Establishment of a new auxin gradient prior to abscission of the lateral root cap. (Manuscript).
- III Philippe Ranocha, Oana Dima, Réka Nagy, Judith Felten, Claire Corratgé-Faillie, Ondřej Novák, Kris Morreel, Benoît Lacombe, Yves Martinez, Stephanie Pfrunder, **Xu Jin**, Jean-Pierre Renou, Jean-Baptiste Thibaud, Karin Ljung, Urs Fischer, Enrico Martinoia, Wout Boerjan and Deborah Goffner (2013). *Arabidopsis* WAT1 is a vacuolar auxin transport facilitator required for auxin homeostasis. *Nature Communications* 4, 2625.

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The contribution of Xu Jin to the papers included in this thesis was as follows:

- I Planning, performance of all the experiments, analysis and summary of the results, preparation and writing part of the manuscript.
- II Planning, performance of all the experiments, analysis and summary of the results, preparation and writing part of the manuscript.
- III Performance of all the experiments for Figure 3 and part of the experiments for Figure 4 and Figure S3.

Abbreviations

1-MCP	1-methyl-cyclopropene
2,4-D	2,4-dichlorophenoxyacetic acid
ACC	1-aminocyclopropane-1-carboxylic acid
ACS	1-AMINOCYCLOPROPANE-1-CARBOXYLATESYNTASE
AGL	AGAMOUS-LIKE
AGP	ARABINOGALACTAN-PROTEIN
ARF	AUXIN RESPONSE FACTOR
ARP	ACTIN-RELATED PROTEIN
ATAF	ARABIDOPSIS TRANSCRIPTION ACTIVATION FACTOR
AUX1	AUXIN RESISTANT1
AVG	amino-ethoxyvinyl glycine
AZ	abscission zone
BC	border cells
BLC	border-like cells
BRN	BEARSKIN
CF	9-hydroxyfluorene-9-carboxylic acid
CTR	CONSTITUTIVE TRIPLE RESPONSE
CUC	CUP-SHAPED COTYLEDON
DAB	DELAYED FLORAL ORGAN ABSCISSION
EIN	ETHYLENE-INSENSITIVE
ERS	ETHYLENE RESPONSE SENSOR
ETR	ETHYLENE RECEPTOR
<i>Gr</i>	<i>Green-ripe</i>
HAE	HAESA
HSL	HAESA-like
IAA	Indole-3-acetic acid
iaaL	IAA-Lys synthetase
iaaM	Trp-2-monoxygenase

<i>ida</i>	<i>inflorescence deficient in abscission</i>
<i>ind</i>	<i>indehiscent</i>
KD1	KNOTTED1-LIKE HOMEODOMAIN PROTEIN1
LAX1	LIKE AUX1
Le	<i>Lycopersicon esculentum</i>
LRC	lateral root cap
LRR-RLK	LEUCINE-RICH REPEATS- RECEPTOR-LIKE KINASE
LX	LX ribonuclease
MKK	MITOGEN-ACTIVATED PROTEIN KINASE KINASE
MPK	MITOGEN-ACTIVATED PROTEIN
MS	Murashige and Skoog
NAA	1-naphthaleneacetic acid
NAC	NAM, ATAF, and CUC transcription factor
NAM	NO APICAL MERISTEM
NPA	1-N-Naphthylphthalamic acid
<i>Nr</i>	<i>Never-ripe</i>
PG	POLYGALACTURONASE
PI	propidium iodide
PIN	PIN-FORMED
PK	Pro transporter
QC	quiescent centre
RNAi	RNA interference
RNase	ribonuclease
SMB	SOMBRERO
TNK4	TOMATO KNOTTED4
WAT	WALLS ARE THIN
WEI	WEAK ETHYLENE INSENSITIVE

1 Introduction

Abscission is a widespread physiological process in plants that plays a variety of roles in development. During their life cycle, plants can shed entire organs and tissues, e. g. leaves, flowers, fruits and the bark in response to developmental or environmental cues (Siwecki & Kozlowski, 1973; Addicott, 1982; Sexton & Roberts, 1982; Osborne & Morgan, 1989; Lewis *et al.*, 2006). As diverse the nature of the shed organs is, as different is the significance or function of separation for plant development. For example, shedding the leaves in autumn minimizes water loss in the winter; separating seeds from fruits permits propagation and rapidly detaching damaged or infected organs is a successful defense strategy of the plant.

Organ abscission usually occurs at a histologically distinct boundary between the organ and the plant body referred to as the abscission zone (AZ). The AZ is comprised of layers of functionally specialized cells with morphologically distinct features like smaller, square-shaped cells, interconnected by branched plasmodesmata and dense cytoplasm (Brown & Addicott, 1950; Gawadi & Avery, 1950; Sexton & Roberts, 1982; Taylor & Whitelaw, 2001). Two types of AZs have been reported, the primary and secondary AZ. Primary AZs differentiate early in development, simultaneously with the development of the lateral organ at a predefined position (Addicott, 1982; Osborne & Morgan, 1989; Taylor & Whitelaw, 2001). By contrast, secondary AZs, also referred to as adventitious AZs, are formed as a response to non-developmental cues, usually late in the development of the lateral organ (Addicott, 1982; Webster & Leopold, 1972; Pierik, 1977, 1980). The number of AZ layers is highly variable between species (Taylor & Whitelaw, 2001; Estornell *et al.*, 2013). While maturation of AZs may be reached early during development, induction of the separation process may occur only in late stages of an organ's life span (Estornell *et al.*, 2013). During the actual separation process cells within the AZ lose cell-to-cell adhesion due to the dissolution of

the pectic matrix in the middle lamella by the secretion of hydrolytic enzymes into the apoplast (Sexton, 1997).

Patterson *et al.* (2001) proposed a general model for abscission processes (reviewed in Estornell *et al.*, 2013), which defines four major steps:

- I Differentiation of the AZ in the future position of organ detachment.
- II Competence acquisition by cells of the AZ to react to abscission signals.
- III Activation of the abscission process within the AZ and organ detachment.
- IV Differentiation of a protective layer.

Although the various abscission processes have common features, upstream signaling, which induces the formation of distinct separation structures and activation of the separation process are thought to be different (Taylor & Whitelaw, 2001). Many lines of research indicate that different plant hormones play important roles in the regulation of AZ formation and activation of abscission. Whereas some hormones act as inhibitors of abscission, like auxin, gibberellins and brassinosteroids (Stutte & Gage, 1990; Ben-Cheikh *et al.*, 1997; Khripach *et al.*, 1999; Taylor & Whitelaw, 2001; Aziz, 2003) others, such as ethylene, abscisic acid, jasmonic acid and cytokinins were shown to accelerate abscission (Sipes & Einset, 1983; Hartmond *et al.*, 2000; Taylor & Whitelaw, 2001; Dal Cin *et al.*, 2007). It is not only the synthesis of these hormones but also their transport and perception, which can be modulated in order to regulate the activity of AZs (Hoad, 1995).

1.1 Leaf abscission

Trees can shed their leaves in response to environmental changes, as seasonal changes in temperature and day-length, or to stress cues, as toxic concentrations of acids or salts, wounding of the leaf blade, pathogen invasion and attack or drought (Taylor & Whitelaw, 2001). Both environmental and stress factors can initiate leaf senescence and as a consequence leaves may generate an abscission-inducing signal, which is transported to the AZ in order to trigger the separation process in primary AZs or to induce the formation of a secondary abscission zone (Addicott, 1982). However, in some cases of stress induced leaf abscission, e.g. drought and pathogen attack, leaf separation can occur in absence of leaf senescence (Taylor & Whitelaw, 2001). Leaf AZs form usually at the base of leaf petioles or leaflets (Addicott, 1982). Simultaneously, or slightly delayed, with the separation phase a protective layer is formed on the proximal site of the fracture. The protective layer reduces water-loss and prevents pathogens from entry (Addicott, 1982; Meirs *et al.*, 2010, 2011; Estornell *et al.* 2013).

1.1.1 Autumnal leaf abscission in deciduous trees

In autumn, plants of many different species drop all of their leaves within short time resulting in seasonally bare, leafless plants. These plants are referred to as deciduous plants. Decreasing day-length and colder temperature in autumn can initiate senescence of leaves in deciduous trees. With the beginning of senescence chlorophyll levels decrease and the photosynthetic capacity of leaves sharply declines (Keskitalo *et al.*, 2005; Fracheboud *et al.*, 2009). Subsequently, anthocyanins accumulate in leaves causing the typical change in colors associated with autumnal senescence (from green to red or yellow; Mol *et al.* 1996; Hoch *et al.*, 2001). Later, the plants remobilize nutrients from the degradation of nucleic acids, proteins and starch in leaves and store them for later use in their roots and stems (Clausen & Apel, 1991; Gan & Amasino, 1997; reviewed in Fischer, 2007).

Comparing with evergreen plants, seasonal leaf abscission in deciduous plants is an evolutionary adaptation to overwinter freezing periods (Zanne *et al.*, 2014). This adaptation allowed deciduous plants to invade vast areas (Freligh & Reich, 2002). In addition, leaf fall can help the plant to generate favorable soil conditions by increasing nutrient release from the soil and regulate the size and composition of microbial communities in the soil (Aponte *et al.*, 2013).

In deciduous trees, e.g. *Populus*, the AZ is formed at the base of the petiole. Secretion of pectinases and cellulases into the apoplast and subsequent dissolution of the middle lamellae are thought to lead to cell separation in the AZ of leaves (Sexton, 1997). After cells of the AZ lose cell-to-cell adhesion, leaves remain attached only through the vascular strands, which are not digested by the secreted hydrolytic enzymes. Completion of abscission by breaking the vascular strands is finally reached by mechanical forces, e.g. wind. The leaf scar, the proximal side of the fracture, is then protected by one or several suberized cell layers.

1.1.2 Retardation of abscission by auxin

In leaf abscission, auxin and ethylene have been pointed out as prominent regulators of abscission (Addicott, 1951; Beyer, 1975). In 1936, La Rue demonstrated that application of auxin onto petioles with excised leaf blades delays petiole abscission in *Coleus* (La Rue, 1936). Later, Addicott reported that the ratio of applied auxin between distal and proximal sides of the AZ is relevant for the timing of abscission and not the absolute concentration of auxin (Addicott *et al.*, 1955; Louie & Addicott 1970). Low auxin concentrations distal to the abscission zone and high concentration proximal favors abscission, whereas the opposite experiment with higher concentrations

on the distal site delays abscission (Louie & Addicott, 1970). Hence, it was suggested that an auxin gradient is spanning the abscission zone and regulates the induction of abscission (Addicott *et al.*, 1955). As a consequence of decreasing auxin concentrations on the distal side either ethylene biosynthesis or/and sensitivity are increased and induce the activity of hydrolytic enzymes degrading cell walls and middle lamellae in the abscission zone (Sexton, 1997).

In Arabidopsis, mutations in two *AUXIN RESPONSE FACTORS* (*ARFs*) have been described to affect petal abscission (Ellis *et al.*, 2005; Okushima *et al.*, 2005). An *arf2* mutation causes delays in the abscission of floral organs, senescence of rosette leaves, flowering time and silique dehiscence, whereas single *arf1* mutants do not display the same phenotypes (Ellis *et al.*, 2005). However, in double mutants of *arf1* and *arf2*, the *arf2* phenotype was strongly enhanced indicating that *ARF1* and *ARF2* have redundant functions in the regulation of abscission and senescence in Arabidopsis (Ellis *et al.*, 2005). Additionally, in *arf2* flowers three members of the *1-AMINOCYCLOPROPANE-1-CARBOXYLATESYNTHASE* (*ACS*) family – *ACS2*, *ACS6* and *ACS8* show decreased transcript levels (Okushima *et al.*, 2005).

In the auxin influx facilitators mutants *aux1* (*auxin resistant1*), *lax1* (*like aux1*) and *lax3* the petal break strength at flower position 3 and 4 is lower than in wild type indicating that the reduction of auxin concentration within the AZ, at the site of their expression, can accelerate petal abscission (Basu *et al.*, 2013). Surprisingly, in an *aux1 lax3* double mutant and an *aux1 lax1 lax2 lax3* quadruple break strength is not more drastically decreased than in the single mutants. In order to test if the manipulation of auxin concentration in the petal AZ affects organ abscission Basu and co-workers expressed two bacterial genes, *iaaM* (Trp-2-monooxygenase) and *iaaL* (IAA-Lys synthetase), under the control of a polygalacturonase, *ADPG2* (*ARABIDOPSIS DEHISCENCE ZONE POLYGALACTURONASE2*), promoter in Arabidopsis (Basu *et al.*, 2013). The *ADPG2* promoter drives gene expression locally restricted to floral abscission zones (González-Carranza *et al.*, 2002). AZ-specific expression of *iaaM*, which converts tryptophan into indole-3-acetamide, causes increased petal break strength. By contrast, expression of *iaaL*, which inactivates IAA, under the control of *pADPG2* decreases petal break strength. These results indicate that the manipulation of auxin concentration in the AZ is sufficient to regulate petal break strength.

Similar as in Addicott's experiments, which led to the auxin gradient hypothesis, distal application of auxin (IAA) to floral explants delays organ separation in tomato (Roberts *et al.*, 1984). Exogenous auxin, however, has no effect on the morphology of the AZ (Roberts *et al.*, 1984). Microarray gene expression analysis from RNA of tomato flower AZs revealed that seven

AUXIN/INDOLE-3-ACETIC ACID (Aux/IAA) genes are down-regulated upon flower removal, whereas three homologs of the IAA-amino acid conjugate hydrolase IAA-LEUCINE RESISTANT (ILR) gene family are up-regulated within two hours after flower removal (Meir *et al.*, 2010). The same authors found for two members of the class I KNOTTED1-LIKE HOMEODOMAIN (KNOX) homeodomain transcription factor gene family, *TOMATO KNOTTED4 (TNK4)* and *KNOTTED1-LIKE HOMEODOMAIN PROTEIN1 (KDI)*, increased expression in the pedicel AZ of tomato (Meir *et al.*, 2010). Ma *et al.* (2015) showed that *KDI* expression is restricted to the AZ of petioles and petals and that overexpression of *KDI* results in accelerated petal abscission. Conversely, when *KDI* is silenced, abscission is delayed, correlating with higher endogenous auxin concentrations and expression of auxin-related genes (Ma *et al.*, 2015).

The above summarized results are consistent with those reported by Hänisch Ten Cate and Bruinsma (1973) who found that although auxin application delays abscission in pedicels of *Begonia* flower buds, it does not affect the anatomy of the abscission zone. Collectively, research on the role of auxin in abscission suggests that auxin can delay cell separation by inhibiting the expression of certain cell wall degrading enzymes (Hong *et al.*, 2000; Tucker *et al.*, 2002), but does not affect AZ differentiation (Ellis *et al.*, 2005).

While direct evidence for auxin or auxin response distribution across the AZ zone in leaf axils still is scarce, Sorefan *et al.* (2009) described a local auxin minimum to be required for the differentiation of a separation zone in the valve margins of the *Arabidopsis* silique. In wild type, the separation layers of the valve margin differentiate at the position of a local auxin response minimum before fruit opening. By contrast, in the *indehiscent (ind)* mutant, which is defective in valve separation, increased auxin response occurs at the position, where normally valve margins differentiate. Similarly, expression of the bacterial auxin biosynthesis gene *iaaM* under the control of the *IND* promoter prevents the differentiation of valve margins (Sorefan *et al.*, 2009). Together, these results support the hypothesis that a local auxin minimum is required for the formation of a functional separation zone.

1.1.3 Acceleration of abscission by ethylene

In 1931, Zimmerman *et al.* reported that high concentration of ethylene caused premature opening and dropping of petals in rose flowers (Zimmerman *et al.*, 1931). Since then, ethylene function in abscission has been intensely researched and various independent lines of evidence show that ethylene is a positive regulator of abscission, which accelerates separation of different organs, such as leaves, flowers and fruits (Brown, 1997; Jackson & Osborne,

1970). During petiole abscission, ethylene concentration in the AZ often increases just before the onset of organ separation (Jackson *et al.*, 1972, 1973). Increasing ethylene concentrations are believed to activate the expression of genes encoding cell wall remodeling enzymes, such as cellulases and pectinases, and their secretion into the apoplast of AZ cells (Addicott, 1982; Brown, 1997).

Genetic evidence for an involvement of ethylene response in abscission came from the study of the Arabidopsis ETHYLENE RECEPTOR1 (ETR1), which contains an amino-terminal domain that possesses ethylene-binding activity (Schaller & Bleecker, 1995). The carboxy-terminal domain of ETR1 exhibits sequence homology to bacterial two-component regulators (Chang *et al.* 1993), which are predominantly sensors and signal transducers of environmental stimuli in a variety of adaptation responses in bacteria (Parkinson & Kofoid, 1992). In the dominant *etr1-1* mutant, which has a missense mutation in the amino-terminal ethylene-binding domain and which is completely insensitive to exogenous ethylene (Chang *et al.* 1993), floral organ abscission is delayed (Bleecker & Patterson, 1997). Similarly, in *ethylene-insensitive2-1* (*ein2-1*), a mutant of the transmembrane protein EIN2 involved in ethylene signaling, floral organ abscission is delayed (Patterson & Bleecker, 1997 and 2004).

In tomato, when floral explants are exposed to atmospheric ethylene, abscission of pedicels is more rapid than in the ethylene-free atmosphere. By contrast, when the explants are pretreated with amino-ethoxyvinyl glycine (AVG), an inhibitor of ethylene biosynthesis, abscission was delayed (Roberts *et al.*, 1984). In line with this, the pretreatment of AZ from tomato explants with an inhibitor of ethylene action, 1-methyl-cyclopropene (1-MCP), causes retardation of abscission, too (Meir *et al.*, 2010).

Significant inhibition of leaf abscission has been observed in LX-deficient (LX ribonuclease (RNase)) transgenic tomato. Expression of the *LX* gene is AZ-specific in tomato and is activated by ethylene in detaching young leaves (Lers *et al.*, 1998 and 2006). In *LX*-deficient antisense lines, both leaf senescence and abscission are delayed (Lers *et al.*, 2006). Furthermore, when the leaves are pretreated with auxin for 30 min and then moved into ethylene containing atmosphere for 48 hours, LX protein levels of auxin pretreated material are lower than the levels of only ethylene-gas treated leaves. This indicates that auxin nullifies the ethylene-induced expression of the LX protein in the AZ of tomato leaves (Bar-Dror *et al.*, 2011).

Three partially ethylene-insensitive, dominant mutants, *Never-ripe* (*Nr*), *Never-ripe 2* (*Nr-2*) and *Green-ripe* (*Gr*), show defects in fruit ripening in tomato. *Nr*, *Nr-2* and *Gr* also affect abscission at the floral pedicel (Wilkinson

et al., 1995; Lashbrook *et al.*, 1998; Barry *et al.*, 2005). *Nr* encodes a homologue of the Arabidopsis *ERS* (*ETHYLENE RESPONSE SENSOR*) and affects the ethylene receptor *LeETR3* as an inhibitor of ethylene signal transduction (Wilkinson *et al.*, 1995; Lashbrook *et al.*, 1998).

The tomato genes *ETHYLENE-INSENSITIVE3* (*EIN3*)-LIKE *LeEIL1*, 2 and 3, homologs of the Arabidopsis *EIN3* that encodes a nuclear-localized component of the ethylene signal-transduction pathway with DNA-binding activity, are known to be involved in the ethylene-mediated initiation of abscission (Tieman *et al.*, 2001). *LeEIL* antisense plants are ethylene insensitive, resulting in flower organ abscission to be delayed until the point when fruit development begins (Tieman *et al.*, 2001). Okabe *et al.* also reported two ethylene intensive mutants of *ETR1* in tomato, *Sletr1-1* and *Sletr1-2*. Both *Sletr1* mutants showed delayed petal abscission (Okabe *et al.*, 2011).

In both, Arabidopsis and tomato, most of the reported ethylene mutants, which show delayed organ abscission, are dominant or semi-dominant. In absence of genetic data from recessive mutants, with the notable exception of *ein2*, this may suggest that although ethylene can modulate the timing of abscission, it may not be essential for the activation of organ abscission (reviewed in Estornell *et al.*, 2013). Therefore and since the inhibition of abscission in completely ethylene-insensitive mutants is incomplete, an ethylene-independent pathway participating in the regulation of organ abscission has been proposed (Bleecker & Patterson, 1997).

In Arabidopsis, a T-DNA mutant of *INFLORESCENCE DEFICIENT IN ABSCISSION* (*IDA*), was the first ethylene-sensitive mutant, with a phenotype in floral organ shedding. The *ida* mutant develops a normal AZ, but is deficient in floral abscission. In *ida*, exogenous ethylene induces senescence of floral organs but is not sufficient for their abscission (Butenko *et al.*, 2003). Since then, the *IDA* signaling pathway in Arabidopsis has been studied intensively. The mutants of the receptor-like protein kinases *HAESA* (*HAE*), the double mutant of *HAESA-LIKE 2* (*HSL2*) and *HAE*, a tandem RNAi transgenic line of *MKK4* and *MKK5* (*MITOGEN-ACTIVATED PROTEIN KINASE KINASE*), and the mutated *MPK6* (*MITOGEN-ACTIVATED PROTEIN KINASE6*) in the *mpk3* background, all have an abscission defective phenotype (Jinn *et al.*, 2000; Cho *et al.*, 2008). Genetic interaction studies have shown that there is a signaling cascade from the putative ligand (*IDA*) to receptors (*HAE* *HSL2*) to cytoplasmic effectors (*MKK4*, *MKK5*, *MPK3*, and *MPK6*) regulating floral organ abscission in Arabidopsis (Stenvik *et al.*, 2008; Cho *et al.*, 2008; Butenko *et al.*, 2012).

Subsequently, the recessive mutants *delayed floral organ abscission 1* (*dab1*), *dab2* and *dab3* were reported to delay floral organ separation (Patterson & Bleecker, 2004). Delayed floral organ detachment in Arabidopsis was also observed in several other mutants and transgenic lines, such as in plants with constitutive expression of *AGL15* (*AGAMOUS-LIKE 15*), a MADS domain transcription factor (Fernandez *et al.*, 2000), and in knock-down lines of *ARP7* (*ACTIN-RELATED PROTEIN7*; Kandasamy *et al.*, 2005). All of the above referred mutants senesce normally and are ethylene-sensitive.

The available evidence clearly shows that ethylene plays an important role during organ abscission but also that ethylene signaling is not a universal requirement or prerequisite for abscission to take place (Addicott, 1982; Taylor & Whitelaw, 2001; Lewis *et al.*, 2006; Estornell *et al.*, 2013). Signaling, which leads to the separation of an organ might therefore not be performed by a single hormonal pathway or a linear sequence of processes (Patterson & Bleecker, 2004).

1.1.4 Cross talk between ethylene and auxin

It is commonly accepted that abscission of diverse organs is regulated by the contrary cooperation between auxin and ethylene action. It is believed that decreased flux of auxin through the AZ triggers increased ethylene signaling and in turn increases the expression and secretion of hydrolytic enzymes involved in the dissolution of the middle lamellae (Roberts & Osborne, 1981; Sexton & Roberts, 1982; Osborne, 1989; Sexton, 1997; Taylor & Whitelaw, 2001). On the other hand, when the flux of auxin through the AZ is high, cell separation is inhibited resulting in organ retention (Addicott, 1982; Sexton *et al.*, 1985). Hence, free auxin in the AZ regulates the sensitivity to ethylene, and therefore any factor, which modulates auxin biosynthesis, homeostasis, signaling or transport in AZ, can also affect the sensitivity of cells in the AZ to ethylene. In order to investigate the cooperation between auxin and ethylene on a molecular and biochemical base, Meir and coworkers examined the abscission-related gene expression in tomato flower AZs after detaching the flowers (Meir *et al.*, 2010). Removal of the flower, the sole auxin source, is expected to deplete the AZ of auxin (Meir *et al.*, 2010). The expression levels of some *AUX/IAAs*, which are directly regulated by auxin, decreases sharply within 2 h after flower removal and remains low thereafter (Meir *et al.*, 2010). As the AZ becomes sensitive to ethylene because of auxin depletion, some ethylene related genes and transcription factors increase their expression levels, such as *ETHYLENE-RESPONSIVE1* (*ER1*), *ETHYLENE RESISTANT4* (*ETR4*), *CONSTITUTIVE TRIPLE RESPONSE1* (*CTR1*), and *ETHYLENE-RESPONSIVE FACTORS* (*ERF1c*). Along with ethylene signaling-related

genes, genes encoding cell wall-modifying proteins are up-regulated (Meir *et al.*, 2010).

Many lines of evidence indicate that ethylene can inhibit auxin transport and biosynthesis, in other words, ethylene can render the AZ more sensitive to itself via positive feedback (Beyer & Morgan, 1971; Beyer, 1973). Beyer and Morgan concluded that following leaf senescence the up-regulation of ethylene signaling reduces auxin transport. Such reduced auxin transport could lead to increased sensitivity to ethylene of the cells in the AZ, and could also induce the production of the enzymes required for abscission and regulate their secretion into the cell wall (Beyer & Morgan, 1971). More complicatedly, auxin itself can stimulate ethylene production and thus accelerate abscission (Abeles & Rubinstein, 1964; Morgan & Hall, 1964).

In Arabidopsis, the inhibition of auxin transport and biosynthesis by ethylene has been studied, too. *WEAK ETHYLENE INSENSITIVE 2* (*WEI2*) and *WEI7* regulate auxin biosynthesis in root tip. *WEI2* and *WEI7* encode subunits of the anthranilate synthase, a rate-limiting enzyme in tryptophan biosynthesis. Overexpression of *WEI2* and *WEI7* results in the accumulation of auxin in the tip of primary root, whereas in the loss-of-function mutants auxin levels are decreased (Stepanova *et al.*, 2005). Subsequently, Swarup *et al.* (2007) reported that IAA biosynthesis in the root tip was inhibited by AVG treatment indicating that ethylene regulates auxin biosynthesis positively in order to enhance the inhibition of root cell elongation. Furthermore, expression of auxin efflux carriers, such as *PIN-FORMED1* (*PIN1*), *PIN2*, *PIN4*, and the influx carrier *AUX1* increases in response to ethylene (Ruzicka *et al.*, 2007). All these results elucidate that ethylene increases the auxin biosynthesis rate and modulates the capacity of polar auxin transport (Ruzicka *et al.*, 2007).

Since Rubinstein and Leopold (1963) proposed a hypothesis of auxin-ethylene interaction during the abscission process, many scientists have modified and improved this model. A recent model for auxin-ethylene crosstalk in leaf and flower abscission is shown in Figure 1 (Taylor & Whitelaw, 2001). Abscission is divided into two phases, the lag and separation phase, and the lag phase is subdivided into stage I and II. During stage I of the lag phase auxin biosynthesis and transport through the AZ are stable, thus the AZ is insensitive to ethylene signaling. In stage II, auxin biosynthesis in the lateral organ and transport across the AZ decline because of environmental and stress factors. The reduction of auxin renders the AZ sensitive to ethylene and allows ethylene signaling in the abscission zone to take place (Rubinstein & Leopold, 1963; Abeles & Rubinstein, 1964; Sexton & Roberts, 1982; Taylor & Whitelaw, 2001). Increasing ethylene signaling induces the secretion of

pectinases and cellulases to hydrolyze the middle lamellae and the cell walls in the AZ (Addicott, 1982; Brown, 1997).

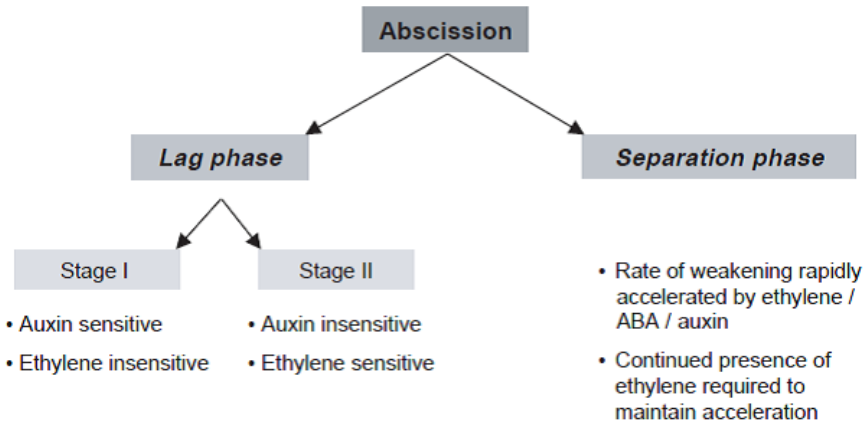


Figure 1. Auxin and ethylene interplay in leaf and flower abscission. Picture reproduced from Taylor and Whitelaw (2001) with the kind permission of the publisher.

1.2 Separation of root border-like cells in Arabidopsis

Like organ separation processes, the separation of the outermost cell layers from the root tip is controlled by phytohormones and environmental factors, and involves the degradation of cell walls and middle lamella (reviewed in Barlow, 2002). Expression of hydrolytic enzymes catalyzing the dissolution of the cell wall and pectin is increased during root cap separation (Driouich *et al.*, 2007).

The root cap plays an important role in the reception of the gravitropic stimuli, in the redirection of growth and in the reduction of mechanical resistance between the soil-root-interface (Hasenstein & Evans, 1988; Fortin & Poff, 1991; Takahashi, 1997; Eapen *et al.*, 2003). The root cap protects its own apical meristem, the so-called root cap meristem, whose initials divide and push their derivatives into direction of the tip. The morphology and the size of the root cap are different in various species. Root caps contain two parts, the columella cells, aligned along the central axis of the root, and the peripheral root cap cells (LRC). Columella cells originate from columella initials, neighboring the quiescent center. LRC cells are derived from the LRC initials at the flanks of the columella initials. Interestingly, epidermal cells of the root share the same initials with the LRC (Fig. 2; Dolan *et al.*, 1993; Wenzel & Rost, 2001; Wenzel *et al.*, 2001; Nawy *et al.*, 2005; Sarkar *et al.*, 2007). Columella cells contain large amyloplasts that serve as statoliths for the

perception of gravity (Moore *et al.*, 1986; Yoder *et al.*, 2001; Saiki and Sato, 2004). The outer two cell layers of the LRC and the tip of the root cap synthesize and secrete the polysaccharide-containing mucilage (Bacic *et al.*, 1986; Morel *et al.*, 1986; Staehelin *et al.*, 1990; Iijima *et al.*, 2004; Cai *et al.*, 2013). The mucilage can lubricate the root so that it penetrates the soil with less mechanical resistance. Prevention of desiccation and facilitating the establishment of mycorrhizae and the interaction with symbiotic bacteria are other functions of the mucilage (McCully & Sealey, 1996; Staehelin *et al.*, 1990; Iijima *et al.*, 2004; Cai *et al.*, 2013).

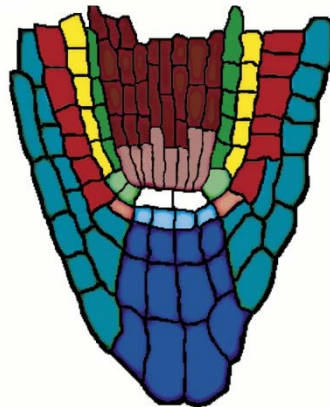


Figure 2. Scheme of the *Arabidopsis* root tip. The primary root meristem consists of the QC (white), and the undifferentiated initial cells of the columella (light blue), the lateral root cap and epidermis (light red), cortex and endodermis (light green). Picture reproduced from Nawy *et al.*, (2005) with the kind permission of the publisher, www.plantcell.org. Copyright American Society of Plant Biologists.

Two types of cells sloughed from the root cap of higher plants have been observed, the border cells (BC; Hawes *et al.*, 2000) and the border-like cells (BLC; Vicré *et al.*, 2005). The BCs are released as dispersed single cells after the mucilage swells due to contact with the soil water (Hawes *et al.*, 1991; Brigham *et al.*, 1998; Hawes *et al.*, 2000). The BCs play an important role to protect the root system against biotic and abiotic stress (Miasaka & Hawes, 2001; Pan, 2004; Gunawardena & Hawes, 2002; Gunawardena, 2005). Unlike the BC, the BLC are released in blocks of associated cells. Despite this, BLCs have the same functions as BCs. The root cap of the model plant *Arabidopsis thaliana* is shed in blocks of associated cells of single layers (Vicré *et al.*, 2005; Hamamoto *et al.*, 2006). The BLC of *Arabidopsis* share characteristic features with BLC/BC from other species, such as they are rich in mitochondria, endoplasmic reticulum, multi-vesicular bodies, Golgi stacks with many Golgi-

derived vesicles, suggesting that these cells actively secrete pectic polysaccharides and arabinogalactan-proteins (AGPs) to their cell walls and surrounding medium (Vicré *et al.*, 2005; Nguema-Ona *et al.*, 2014). While the traditional model species to study BC function are cotton (*Gossypium hirsutum*), wheat (*Triticum aestivum*), maize (*Zea mays*), tomato (*Lycopersicon esculentum*) and pea (*Pisum sativum*), research on BLC can be performed in *Arabidopsis* (Vicré *et al.*, 2005).

The separation of BLC in *Arabidopsis* starts at the age of 5 to 7 days. Before, the root cap stays intact (Vicré *et al.*, 2005). Lateral root cap layers are shed one by one, starting from the outermost, oldest layer. In 13-days-old seedlings, at least 3 layers of BLC are shed (Vicré *et al.*, 2005). In *Arabidopsis*, separation of the BLC occurs before cell death and BLC remain viable for at least 24 hours after separation (Vicré *et al.*, 2005). Similarly, BC also can survive after the separation from the root cap and even survive *in vitro* in similar conditions as in nature. Interestingly, BC can divide and develop into callus tissue *in vitro* (Hawes *et al.*, 2000; Hawes & Lin, 1990).

The separation of BC in pea (*Pisum sativum*) revealed that a specific polygalacturonase enzyme, which can hydrolyze polygalacturonic acid to disassemble primary cell walls, is activated during the BC separation process (Hawes & Lin, 1990). In comparison with BC, BLC in *Arabidopsis* secrete significant amounts of homogalacturonan and AGPs into their cell walls so that they do not become dispersed individually into suspension. The *Arabidopsis root epidermal bulger1-1 (reb1-1)* mutant, which has reduced levels of AGPs, releases less BLC with altered morphology compared to wild-type roots (Driouich *et al.*, 2007). This has been interpreted in such a way that there is a specific cell wall composition or structure in the BLC, that confers resistance to hydrolysis of cell walls or that the BLC have no functional pectolytic enzymes capable to hydrolyze the pectic matrix of cell walls (Driouich *et al.*, 2007).

So far research on the separation processes of the BC and BLC has more focused on regulatory functions of the cell wall modifying enzymes than on the effects of phytohormones. For the BC separation in maize, Ponce demonstrated that the interruption of polar auxin transport by the application of the polar auxin transport inhibitor NPA (1-N-naphthylphthalamic acid) increased the BC shedding (Ponce *et al.*, 2005). Surprisingly, application of ACC (1-aminocyclopropane-1-carboxylic acid) and of the ethylene synthesis inhibitor AVG (aminovinylglycin) decreased the BC shedding significantly (Ponce *et al.*, 2005). These results indicate that the BC separation is regulated by ethylene and by auxin, and that auxin and ethylene may act coordinately to regulate BC differentiation and separation, in a similar fashion as in other

abscission processes (Ponce *et al.*, 2005; Driouich *et al.*, 2007). For the BLC separation in *Arabidopsis*, no such data is available. But for a mutant of the auxin efflux carrier *PIN4*, an increased number of columella layers has been reported (Friml *et al.*, 2002).

In *Arabidopsis*, the NAC domain transcription factors, *SOMBRERO* (*SMB*), *BEARSKIN1* (*BRN1*), and *BRN2*, are required for the BLC separation (Bennett *et al.*, 2010). The *Arabidopsis* genome contains more than 100 members of the NAC domain transcription factors (Olsen *et al.*, 2005) and many of them have been reported to be regulated by phytohormones (He *et al.*, 2005). *SMB*, *BRN1* and *BRN2* regulate independently and redundantly the differentiation of root cap cells, including their morphology and their ability to separate from the root. In the *smb-3* mutant, the LRC cells display an abnormal morphology and fail to detach from the root tip. In the *brn1-1 brn2-1* double mutant, the separation of BLC is defective, but the shape of the BLC was normal. Surprisingly, in the *smb-3 brn1-1 brn2-1* triple mutant both the shape and the detachment of the BLC are altered (Bennett *et al.*, 2010).

2 Objectives

The main aim of the work described in this thesis was to understand how auxin regulates leaf and root cap abscission.

2.1 Leaf abscission in *Populus* (Manuscript I)

The specific objectives of this project were:

- the identification of auxin carriers involved in leaf abscission.
- the examination of the function of auxin and ethylene during leaf abscission.
- the study of auxin - ethylene crosstalk in the abscission zone.

2.2 Root cap abscission in *Arabidopsis* (Manuscript II)

The specific objectives of this project were:

- the study of auxin response during root cap abscission.
- testing the plausibility of a regulatory function of an auxin gradient in root cap abscission.
- the examination of cell wall remodeling during root cap abscission.

2.3 *WAT1* function in plant development and auxin homeostasis (Manuscript III)

In Manuscript I, a *Populus* homolog of the *Arabidopsis* WALLS ARE THIN1 (*WAT1*), a vacuolar auxin transport facilitator, was identified as being regulated during leaf abscission in a microarray experiment. We wanted to study:

- the expression pattern of *WAT1* in *Arabidopsis*.
- the subcellular localization of *WAT1*.
- the function of *WAT1* in development.

3 Materials and methods

3.1 *Populus* as a model tree species to study leaf abscission

The genus *Populus* consists of deciduous flowering tree species, including poplar, aspen, and cottonwood species. In most cases, the flowers are dioicous and the predominant pollination syndrome is anemophily (wind pollination). Natural *Populus* populations occur in temperate climate regions of the northern hemisphere, where these trees shed their leaves in autumn (autumnal leaf abscission). Many *Populus* species are considered to be fast growing with straight, massive stems and can be easily clonally propagated. In 1986, Parsons *et al.* produced the first transgenic woody plants by transforming *P. trichocarpa* × *deltoids* (Parsons *et al.*, 1986). Soon thereafter the transformation protocol had been optimized and other *Populus* hybrids became transformable, such as *P. alba* L. × *P. tremula* (referred to as *Populus* × *canescens*; Devillard, 1992; Leple *et al.*, 1992) and *P. tremula* × *P. tremuloides* (*Ptt*, referred to as T89; Nilsson *et al.*, 1992).

In 2006, a complete sequence of the *Populus trichocarpa* genome, the first completed tree genome, was published by the Joint Genome Institute together with researchers at the Umeå Plant Science Centre (Tuskan *et al.*, 2006). *Populus trichocarpa* has a comparatively small genome of about 535 mega base pairs. The availability of a full genome sequence facilitated molecular breeding approaches, as genome-wide association mapping, to exploit the huge natural variation within more than 30 *Populus* species (Porth *et al.*, 2013).

After the publication of the *P. trichocarpa* genome (Tuskan *et al.*, 2006), the importance of *Populus sp.* as a model for various aspects of tree physiology has steadily increased. Nowadays, *Populus sp.* is used as a model species for e.g., wood formation, dormancy, drought stress, adventitious root formation, organic volatile emissions and mycorrhiza (Jansson & Douglas, 2007; Fischer & Polle, 2011; Legué *et al.*, 2014; Douglas & Polle, 2010; Ditengou *et al.*,

2015). Rapid growth and easy and efficient clonal mass propagation makes *Populus* interesting as a possible sustainable feedstock for bioenergy production (Porth & El-Kassaby, 2015).

In Manuscript I, two different *Populus* hybrids were employed to study the regulation of leaf abscission, i.e. hybrid aspen T89 and *Populus* × *canescens*.

3.2 The Arabidopsis root as a model to study root-cap abscission

Arabidopsis is a popular model organism for most aspects of plant biology. Despite being a complex multicellular eukaryote, *Arabidopsis thaliana* has a relatively small genome of approximately 135 mega base pairs. The small size of its genome made Arabidopsis attractive since the beginning of plant molecular biology. In fact, the Arabidopsis genome was the first plant genome to be fully sequenced in 2000 by the ‘Arabidopsis Genome Initiative’. In addition, Arabidopsis provides many other advantages, such as its short life cycle, prolific seed production, easy cultivation, highly efficient and simple transformation protocols with *A. tumefaciens* and a large number of publicly available mutant lines and genetic resources.

In Manuscript II, we made use of some of the benefits Arabidopsis provides in order to study root cap abscission. Arabidopsis root tips display a simple and for plant organs highly ordered cellular organization. Aspects of patterning, cell specification, growth and differentiation, meristem maintenance, polarity, lateral root formation and physiological and environmental responses have been studied in Arabidopsis in great detail during the last 25 years (Benfey *et al.*, 2010). Arabidopsis roots can grow on solid growth media in transparent petri dishes, which facilitates the visual observation of mutant phenotypes or growth responses. Due to their small diameter Arabidopsis roots are fully transparent in confocal and light microscopy and various protocols have been established to exploit the benefits of live imaging (Petricka *et al.*, 2012).

In Arabidopsis, the root cap consists of the lateral root cap (LRC) and columella. The root cap forms protective layers of cells, which are continuously separated from the root tip (Driouich *et al.*, 2007). The root cap is involved in the reception of gravity and regulates the gravitropic response (Sato *et al.*, 2014). The investigation of lateral root cap cells, the so called border like cells (BLC), that are released to the surrounding media has been pioneered by Vicré *et al.* in 2005. Examination of the morphology, cell wall composition of BLC and of bacteria attachment to BLC in Arabidopsis seedlings has been reported (Driouich *et al.*, 2007).

In Manuscript II, we employed *Arabidopsis* in order to study cell separation in the root cap. All the transgenics and mutants used in Manuscript II are in the *Col-0* background.

3.3 Dark induction of leaf abscission in *Populus*

We adopted a method by covering leaf blades of *Populus* with aluminum bags in order to induce rapid and synchronous leaf abscission in large numbers (Manuscript I). The trees were grown to a height of approximately 2 m and then leaves with a petiolar angle bigger than 75 degrees were shaded with aluminum bags under standard greenhouse conditions (Fig. 3). Only healthy leaves of similar size were considered for the induction experiments. Each aluminum bag was labeled with a unique code referring to the genotype, tree replicate, leaf number and treatment. Trees were gently shaken once per day, the dropped bags collected and the identifiers recorded.



Figure 3. Dark induced leaf abscission in *Populus*. Fully expanded leaves were covered by aluminum bags.

3.4 Gene expression analysis

Fully expanded leaf blades of hybrid aspen, *Populus tremula* L. × *P. tremuloides* Michx.; clone T89, were shaded in aluminum foil and total RNA was extracted from 3-mm-thick leaf axils, 6 days after shading started, using RNeasy Plant Mini Kit (Qiagen). 2 µg of total RNA was used as a template for reverse transcription with the QuantiTect Reverse Transcription Kit (Qiagen). Quantitative real-time PCR was performed using SYBR Green I Master in combination with a Light Cycler 4800 (Roche Diagnostics) qPCR machine.

Primers are specified in Table 1. Expression was normalized to *PtACTIN1* expression. The primers for the qPCR are shown in Table 1.

Table 1. *Primers used for qPCR*

Gene model	Sequence
<i>PtIDA</i>	GACCGGGCGCTACTCTAATGG TGGATGGACCAGAAGGTGGAA
<i>PtIDL1</i>	TGTTCACCTTCGGCTCCAT CCTTGGTGGCCATGTCTG
<i>Pt019G078400</i>	GGGAAGCTGCCTGAGGGAAT GTGAGAACCGGCCGAAATTG
<i>Pt007G135400</i>	CTTTGCCGCTGACCAATCT CGGACAGTGACTCGGGAAGG
<i>PtACTIN1</i>	CGATGCCGAGGATATTCAAC ACCAGTGTGTCTTGGTCTACCC

4 Results and discussion

In Manuscript I, dark-induced leaf abscission in *Populus* was described. We studied the auxin response during abscission and identified auxin transporters, whose expression was regulated during leaf abscission in intact trees. Expression patterns of these auxin transporters were examined and it was also shown that regulation of their expression is independent of ethylene signaling. In the following chapters, additional data in support of the findings in Manuscript I are presented.

4.1 Application of auxin delayed the formation and maturation of the AZ (Manuscript I)

In Manuscript I, we described the expression pattern of the auxin response reporter *GH3::GUS* during dark-induced leaf abscission (Figure 2 in Manuscript I). The AUXIN RESPONSE ELEMENTS (AuxRE) of this promoter had previously been used to construct a synthetic promoter (*DR5*) with better auxin inducibility and lower background expression (Ulmasov *et al.*, 1997). We found a similar expression pattern for the *DR5::GUS* reporter line in T89 than for the *GH3::GUS* reporter. In *DR5::GUS* T89 lines, a new auxin response maximum emerged at the lower (abaxial) side of the petiole after 3 days of shading (Fig. 4A, B). Browning of the AZ indicated the presence of a mature AZ and *DR5::GUS* became weaker in the AZ 9 days after shading started (Fig. 4C, D). Local application of the synthetic auxin 2,4-D (100 μ M 2,4-Dichlorophenoxyacetic acid) to the axils of intact leaves delayed the formation of a new auxin response maximum, which was barely visible at the lower side of the petiole 3 days after dark-induction (Fig. 4E, F). Nine days after shading started, the auxin response was strong in the AZ and proximal to the AZ, but unlike in the mock-treated axils, the cells in AZ did not brown (Fig. 4G, H). These observations are in line with the results presented in

Manuscript I (compare with Figure 4 in Manuscript I), where local application of 2,4-D to the axil of shaded leaves delayed separation by approximately 4 days. Additionally, the results presented here suggest that auxin acts on the formation of the abscission zone.

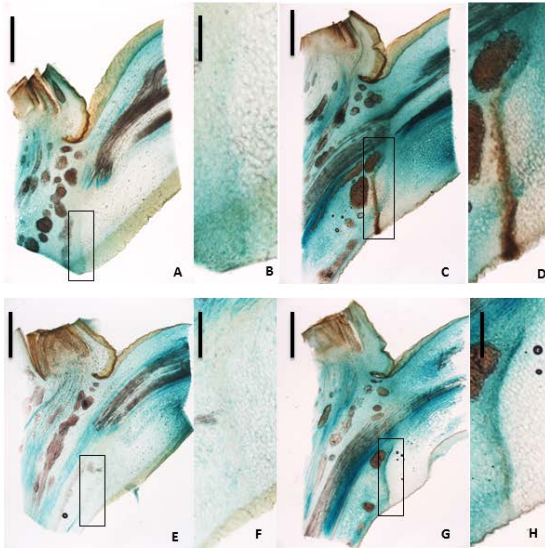


Figure 4. Exogenous auxin delayed the formation of a local auxin response maximum and the maturation of the leaf AZ. Expression of *DR5::GUS* in the AZ of 2,4-D-treated and control transgenic plants. A) to D), dark induction without auxin treatment; A) and B), 3 days after shading started, black frame highlights the AZ; C) and D), 9 days after shading started. Mock treatment, lanolin paste locally applied to the AZ. E) to H), dark induction with application of 2,4-D; E) and F), 3 days after shading started; G) and H), 9 days after shading started. Lanolin paste containing 100 μ M 2,4-D was glued to the leaf axils locally. Scale bars, 1 mm (A, C, E, G); 200 μ m (B, D, F, H).

4.2 Ethylene-auxin crosstalk (Manuscript I)

Auxin and ethylene have antagonistic functions during the abscission of various organs (Hong *et al.*, 2000). To test if auxin acts upstream of ethylene on leaf abscission, we transformed *DR5::eto3* and *DR5::etr1-1* into T89. *eto3* is a dominant mutant of the ethylene biosynthetic gene 1-AMINOCYCLOPROPANE-1-CARBOXYLATE SYNTHASE 9 (*ACS9*) and can induce higher concentrations of ethylene in *Arabidopsis* (Chae *et al.*, 2003). By contrast, the mutant of the dominant *etr1-1* allele in *Arabidopsis* is insensitive to ethylene (Bleecker *et al.*,

1988; Guzmán & Ecker, 1990; Chang *et al.*, 1993; Chen & Bleecker, 1995). For each construct, we isolated two lines with high levels and one line with low levels of transgene expression (data not shown). Leaves of these lines were subjected to induction of leaf abscission by shading under standard greenhouse conditions. For each line, 3 trees and 15 mature leaves for each tree were shaded to analyze timing of leaf abscission. For both constructs, no correlation between transgene expression levels and timing of abscission could be observed (Fig. 5). Assuming that the *DR5* promoter activity was strong enough to modulate either ethylene biosynthesis or signaling, this indicates that the cells with *DR5* promoter activity at the new auxin maximum are not sensitive to overproduction of ethylene and that these cells do not require ethylene signaling to exert their function in leaf abscission. This is in line with the hypothesis formulated in Manuscript I of independent, parallel action of ethylene and auxin on leaf abscission.

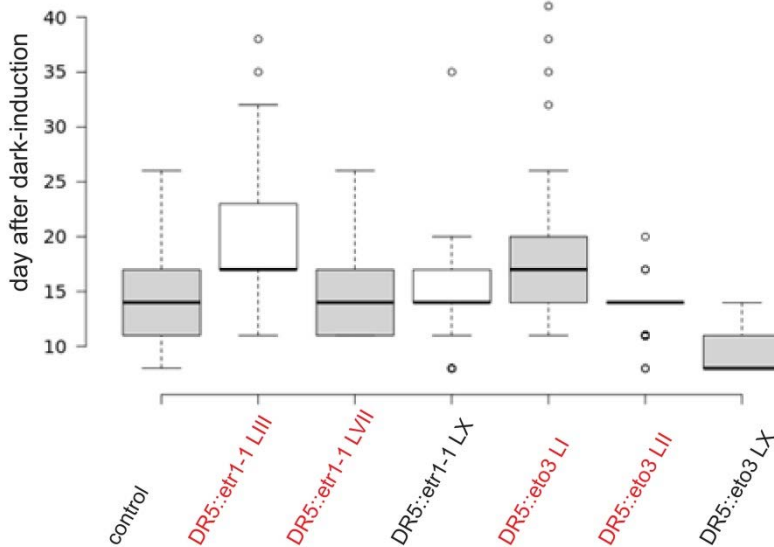


Figure 5. Auxin acts independently of ethylene on leaf abscission. Days after dark-induction started are shown on the y-axis. Box plot, center lines show the medians; box limits indicate the 25th and 75th percentiles; whiskers extend to 5th and 95th percentiles, outliers are represented by dots. 90 individual leaves from 6 T89 trees were shaded as controls. *DR5::etr1-1* lines III and VII had relatively high levels of *etr1-1* expression (red); line X had low levels of transgene expression. *DR5::eto3* line I and II had relatively stronger expression levels of *eto3* (red); line X had low expression levels of the transgene.

4.3 Expression of auxin efflux carriers during floral organ abscission in *Arabidopsis* (Manuscript I)

In Manuscript I, we found that expression of some auxin transporters was down-regulated and spatially restricted during abscission based on the results of qRT-PCR, microarrays and *promoter::GUS* reporter constructs. The very same constructs, *PttPIN1b::GUS*, *PttPIN5b::GUS* and *PttWAT1::GUS*, were transformed into *Arabidopsis*, which does not shed its leaves (Patterson, 2001). For each construct at least 5 independent lines were isolated. All the lines showed intense GUS staining in lateral floral organs and especially strong expression was detected in the protective layer of the abscission zone after the petals and stamen were shed (Fig. 6). This may indicate that factors regulating the expression of *PttPIN1b::GUS*, *PttPIN5b::GUS* and *PttWAT1::GUS* during leaf abscission in *Populus* are conserved in *Arabidopsis*. Thus, regulation of auxin transport during abscission may be conserved in abscission of diverse organs across a wide range of species.



Figure 6. GUS expression in *Arabidopsis* floral AZs. Representative lines for *PttPIN1b::GUS* (A), *PttPIN5b::GUS* (B) and *PttWAT1::GUS* (C) are shown.

4.4 Increased *PtIDA* expression levels in the AZ during leaf abscission (Manuscript I)

In *Arabidopsis*, the *INFLORESCENCE DEFICIENT IN ABSCISSION* (*IDA*) gene was found to regulate the abscission of sepals, petals and stamens. *IDA* encodes a small secreted peptide ligand, which acts upstream of the two leucine-rich repeat receptor-like kinases HAESA (HAE) and HAESA-LIKE2

(HSL2) (Stenvik *et al.*, 2006). Overexpression of *IDA* accelerates the abscission of floral organs (Stenvik *et al.*, 2006), whereas in an *ida* mutant abscission of floral organs is completely impaired (Butenko *et al.*, 2003). Intriguingly, *IDA* expression is inducible by exogenous auxin and its temporal regulation of gene expression is not dependent on ethylene signaling (Kumpf *et al.*, 2013; Butenko *et al.*, 2006). We tested if the expression of the two closest *Populus IDA* homologs, *PtIDA* and *PtIDL1*, are up-regulated during leaf abscission in a similar way as *IDA* expression is induced during floral organ abscission in *Arabidopsis* (Butenko *et al.*, 2003). Both *PtIDA* and *PtIDL1* were significantly up-regulated in dark-induced leaf axils compared to controls (Fig.7). By contrast, expression of the closest *Populus* homologs of *HAE* and *HSL2*, the putative *IDA* receptors, was not affected by the induction of leaf abscission (Fig. 7). Together, these results indicate that the partially ethylene-independent *IDA* signaling plays a common role in the abscission of various organs in different species.

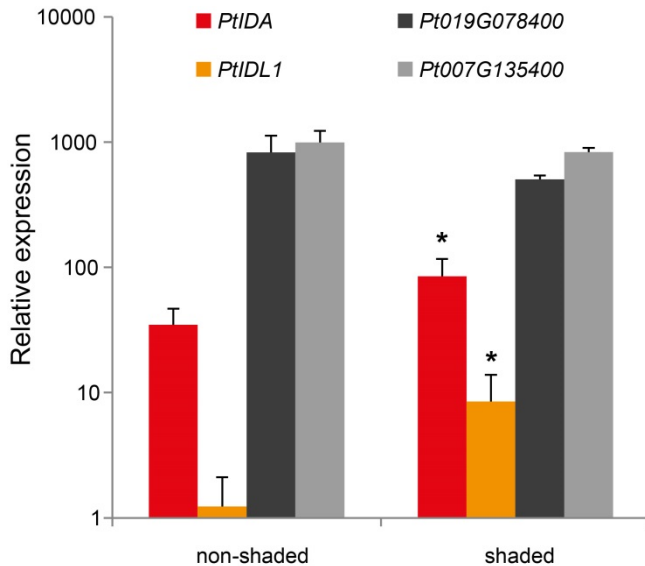


Figure 7. Expression of *PtIDA* and *PtIDL1* in leaf axils is induced upon shading of the leaf blade. Fully expanded leaves were shaded in aluminum foil and gene expression was determined 6 days after shading treatment started. Fifty percent of the shaded leaves were abscised 8 days after the shading treatment started. By contrast, non-shaded leaves were not dropped. Histograms represent gene expression levels normalized versus *PtACTIN1* expression levels. Mean \pm s.d., n=3 biological replicates. *, p < 0.05 by two-tail t-test of non-shaded versus shaded.

4.5 Pectin remodeling during leaf abscission (Manuscript II)

During abscission, cells of the AZ (separation zone) lose cell-to-cell adhesion because of the dissolution of pectin in the middle lamellae and cell walls by increased secretion of pectinases, cellulases and other hydrolytic enzymes (Sexton, 1997). De-esterification is a typical pectin modification in separating cells (Lee *et al.*, 2008). In Manuscript II, we described the distribution of the JIM5 epitope in root cap abscission. The monoclonal antibody JIM5 recognizes partially methylesterified or de-methylesterified homogalacturonan (HG). In *Populus*, prior to separation, JIM5 signals were evenly distributed across the whole tissue, whereas in separating tissues JIM5 fluorescence was largely restricted to the separation zone and tissue proximal to it (Fig. 8). This result suggests that during leaf abscission in *Populus* similar events of cell wall remodeling take place as previously reported for organ abscission in other species.

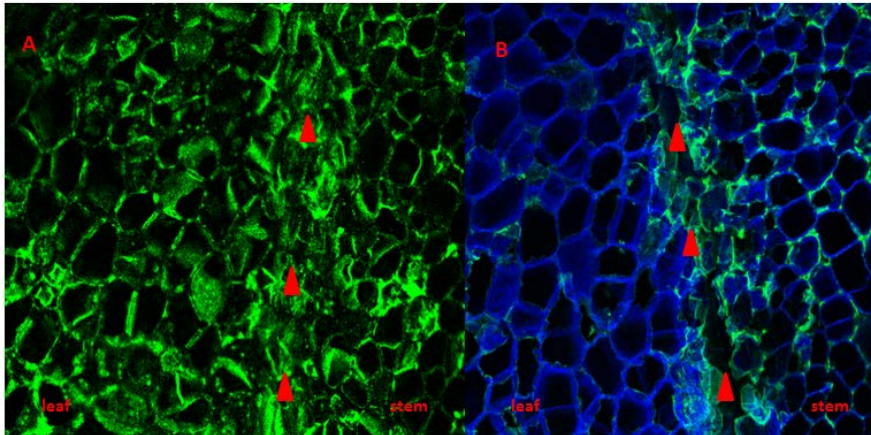


Figure 8. Immunolocalization of the JIM5 epitope recognizing de-methylesterified HG, (green) in the leaf axil of T89. Red arrowheads point to the AZ in A) an axil with a firmly attached petiole and B) an axil where the leaf is separating. Blue in B) shows calcofluor-white stained cell walls.

4.6 Expression of *WAT1* in vascular tissue but not in lateral root caps (Manuscript III)

Auxin plays a crucial role in multiple aspects of plant growth and development; therefore auxin biosynthesis, metabolism, transport and signaling have been intensely studied in the last decades (Benjamins & Scheres, 2008). We have examined the function of auxin in organ separation and although this is a very specific process it is likely that many components of auxin physiology

are not specific for organ abscission but are common players in other auxin-regulated processes. One of the most significantly down-regulated genes during leaf abscission was *PtWAT1* (Manuscript I). At the time we obtained this result, the biochemical function of WAT1 was unknown. However, coregulation with auxin regulated genes and the predicted transmembrane structures suggested that WAT1 could be a so far unknown auxin transporter. In order to better understand the WAT1 function, we collaborated with the Goffner group, which had previously described the *wat1* loss-of-function phenotype (Ranocha *et al.*, 2010). *wat1* mutants are defective in secondary cell wall deposition in interfascicular fibers and xylem vessels and fibers (Ranocha *et al.*, 2010). We localized WAT1-GFP to the tonoplast and our collaborators found that WAT1 transports auxin from the vacuole across the tonoplast to the cytoplasm (Ranocha *et al.*, 2013, Manuscript III). We could rescue the cell wall phenotype by local application of synthetic auxins (Ranocha *et al.*, 2013, Manuscript III). This strongly suggests that auxin is a WAT1 transport substrate in planta.

Next, we came up with the question if *WAT1* is involved in root cap abscission. To this end, we analyzed a *WAT1::GUS* line for root expression. *WAT1::GUS* expression was restricted to the vasculature of the root but not expressed in the lateral root cap (Fig. 9). Neither *wat1* mutants nor the *WAT1* overexpressor lines displayed a phenotype in root morphology or auxin sensitivity of 5-day-old roots (Fig. 10) and there was no obvious abscission related phenotype in 10-day-old roots of *wat1* (Fig. 11). Together, these results indicate that WAT1-mediated auxin homeostasis does not play a significant role in root cap abscission. Alternatively, *WAT1* function could be masked by redundancy. Interestingly, the closest *WAT1* paralog *AT3G53210* is expressed specifically in the lateral root cap and the underlying epidermis (Fig. 12).

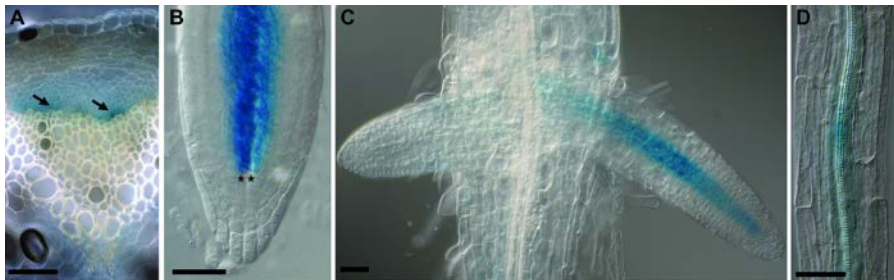


Figure 9. *WAT1::GUS* is expressed in the vasculature and associated tissues but not in the lateral root cap. A), Fascicular bundle. Arrows point to cells differentiating into xylary fibers. Scale bar 100 μm . B), Meristem of a 5-day-old root. GUS activity was detected in stele but not in quiescent center (*) or lateral root cap. Scale bar, 50 μm . C), Lateral roots of a 7-day-old seedling. Before vascularization young lateral root primordia did not express *WAT1::GUS*. Scale bar, 100 μm . D),

WATI::GUS activity was associated with vascular tissue in the elongation zone of the root. Scale bar, 50 μm .

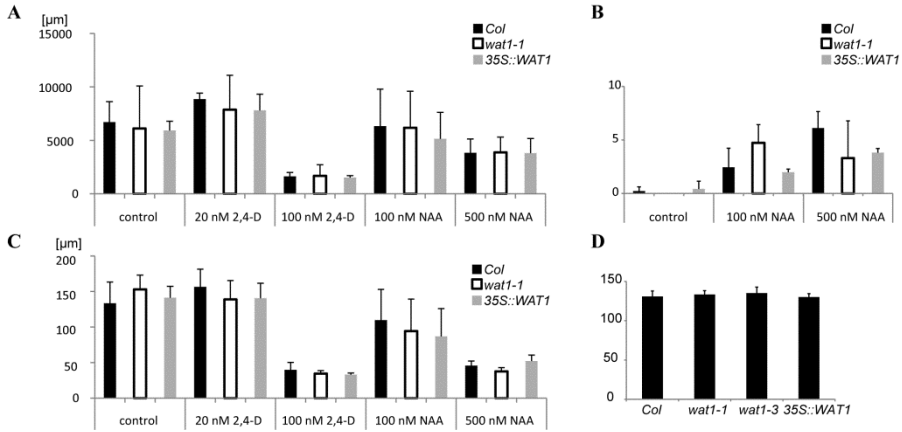


Figure 10. No significant changes in root morphology and auxin sensitivity in 5-day-old *wat1* seedlings. Mean values of three experimental replicates. For each replicate, either 3-5 individuals (A-C) or at least 6 seedlings (D) were used. t-test, $p > 0.05$. A), Root length; B), Number of lateral root primordia; C), Trichoblast length, five trichoblasts per seedling were measured; D), Reorientation angle in gravitropic response, 180° corresponds to no reorientation, 90° complete reorientation in direction of the new gravity vector. Four-day-old seedlings grown on vertical plates were turned by 90° and reorientation angles were measured one day after turning.

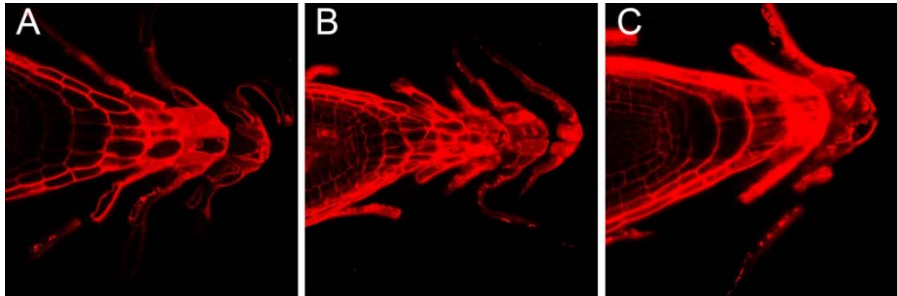
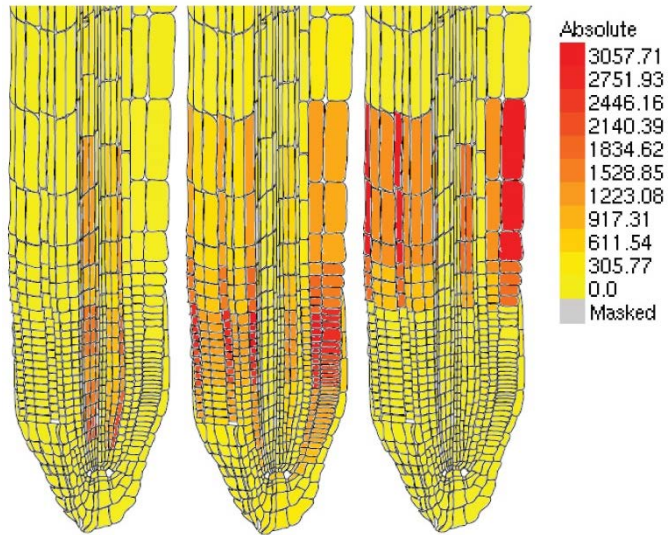


Figure 11. The *wat1-3* mutant does not have any phenotype related to root cap abscission. (A-C), 10-day-old root tips. A), wild type; B), *wat1-3*, and C), *35S::WATI::GFP*.



AT1G75500 AT3G53210 AT3G18200

Figure 12. Lateral root cap specific expression of the closest *WAT1* (*At1G75500*) paralogs. Data derived from Brady et al., 2007 and displayed with the Arabidopsis eFP Browser.

5 Conclusion and future perspectives

Although the season when leaves separate from deciduous trees – “the fall” – is named after “the fall of the leaf” little attention has been paid to how seasonal leaf abscission is regulated. Current models of leaf abscission almost uniquely derive from the study of annual plants. These models, reproduced in popular plant biology text books (e.g. Taiz and Zeiger), suggest that reduced auxin flow from the leaf blade into the leaf axil leads to increased ethylene signaling and consequently to hydrolysis of middle lamellae.

In Manuscript I, we described the establishment of an experimental model for leaf abscission in *Populus* trees. We found that prior to the formation of an abscission zone, a new auxin maximum is established, which likely provides positional cues for the formation of an abscission zone. Inhibition of polar auxin transport, as well as exogenous auxin application, delays abscission. In contrast to the current text book opinion, auxin acts independently of ethylene signaling on leaf abscission.

We identified auxin transport facilitators, which are among the most significantly regulated genes during leaf abscission. We described their expression patterns during leaf abscission (Manuscript I). Functional analysis of these auxin transport facilitators and the determination of their subcellular localization will shed light on how the local auxin maximum is established in the leaf axil.

Unlike in leaf abscission, separation of the root cap does not involve the formation of a morphologically complex abscission zone. This makes the root cap a more accessible model than leaf axils in order to study the basic principles of cell separation. We found that root cap cell layers at the minimum of the columella-spanning auxin gradient undergo abscission. Transport of auxin from its source in the quiescent center to the periphery is hindered due to the absence of auxin efflux carrier expression in the outer layers of the columella (Manuscript II). Solely based on physiological experiments, Addicott (1955) proposed an auxin gradient to regulate organ abscission. Here,

we provide novel, molecular evidence, which is in line with Addicott's hypothesis. Future experiments should include the genetic analysis of auxin gradient formation in the root cap, e.g. by the means of a mutant screen or inducible silencing of PIN expression.

One of the most significantly regulated genes during abscission is a *Populus* homolog of the *Arabidopsis* *WAT1*. At the time of our gene expression study, the biochemical function of *WAT1* was still unknown. In collaboration with other groups, we found that *WAT1* is a tonoplast-localized auxin transporter regulating cellular auxin homeostasis (Manuscript III). *WAT1*, which belongs to a large gene family of transmembrane proteins, is not expressed in the root cap and the perturbation of this gene does not lead to any phenotype related to abscission. Future work should address expression patterns and functions of the close *WAT1* homologs in order to test if auxin homeostasis mediated by *WAT1* homologs plays any role in organ abscission.

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