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Review

Aquaporins and plant transpiration

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ABSTRACT

Although transpiration and aquaporins have long been identified as two key components influencing plant water status, it is only recently that their relations have been investigated in detail. The present review first examines the various facets of aquaporin function in stomatal guard cells and shows that it involves transport of water but also of other molecules such as carbon dioxide and hydrogen peroxide. At the whole plant level, changes in tissue hydraulics mediated by root and shoot aquaporins can indirectly impact plant transpiration. Recent studies also point to a feedback effect of transpiration on aquaporin function. These mechanisms may contribute to the difference between isohydric and anisohydric stomatal regulation of leaf water status. The contribution of aquaporins to transpiration control goes far beyond the issue of water transport during stomatal movements and involves emerging cellular and long-distance signalling mechanisms which ultimately act on plant growth.

Key-words: guard cell signalling; isohydric and anisohydric plants; stomatal movement.

INTRODUCTION

Gas exchange between plant shoots and the atmosphere plays a key role in plant function and performance through the intake of carbon dioxide (CO₂) to supply photosynthetic carbon fixation and the diffusion of water vapour by transpiration. The latter is driven by the evaporative demand, that is the dramatic drop in vapour pressure between plant tissues and the atmosphere. Water loss by transpiration cannot simply be considered as the detrimental component of a trade-off for optimizing CO₂ intake and carbon fixation. Transpiration is important in leaf cooling and drives xylem-mediated mass flow of nutrients from the soil into the uppermost parts of the plant (Cowan & Farguhar, 1977; Taiz & Zeiger, 1991; Medina & Gilbert, 2016). Because of these multiple and sometimes conflicting functions, gas exchanges by plant aerial parts must be tightly controlled. Gas exchanges are mediated to a large extent through stomata, specialized pores differentiated from epidermal cells at the surface of leaves and stems (Taiz & Zeiger, 1991; Murata et al., 2015). The remaining outer surface of the epidermis is covered with a largely gas-tight cuticle (Yeats & Rose, 2013).

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Aquaporins are intrinsic membrane proteins present in the plasma membrane and most inner membranes of plants cells. While most aquaporins function as channels to facilitate transmembrane water transport, they can also transport small neutral molecules such as gases (CO₂, ammonia), reactive oxygen species (hydrogen peroxide: H₂O₂) and metalloids (boric acid, silicic acid, antimonite, arsenite) (Kaldenhoff, 2012; Bienert & Chaumont, 2014; Maurel et al., 2015). The water transport function of aquaporins plays a key role in as diverse processes as root water uptake, leaf hydraulics, seed and pollen grain germination, expansive growth and lateral root emergence (For recent reviews, see Chaumont & Tyerman, 2014; Maurel et al., 2015). In relation to their multiple cellular localizations and functions, plant aquaporins fall into at least four subclasses that each shows a high isoform multiplicity. Of specific interest for this article, are the Plasma membrane Intrinsic Proteins (PIP; 13 isoforms in Arabidopsis) and the Tonoplast Intrinsic Proteins (TIP; 9 isoforms in Arabidopsis) which represent the most abundant aquaporins in the plasma membrane and vacuolar membrane (tonoplast), respectively.

This review examines the relations between aquaporins and plant transpiration, two important components of plant water relations. We first discuss the various facets of aquaporin function in stomatal guard cells and show that it involves transport of water but also of other molecules. We also show that function of aquaporins in roots and shoots is intimately linked to transpiration. Thus, the contribution of aquaporins to transpiration control goes far beyond the issue of water transport during stomatal movements and involves emerging cellular and long-distance signalling mechanisms.

THE ROLE OF AQUAPORINS IN GUARD CELLS

Principles of stomatal movements

Stomata are microscopic pores delineated by a pair of guard cells. In certain plant families such as cereals, guard cells are themselves surrounded by specialized subsidiary cells to form so-called stomatal complexes. Stomatal aperture is itself determined by the volume and mechanics of guard cells. An increase in guard cell osmotic pressure leads to water intake, guard cell swelling and stomatal opening. Conversely, osmotically driven water efflux from guard cells leads to stomatal closure. These movements are under tight control by the circadian clock and numerous environmental and hormonal signals that promote stomatal opening (light, high air humidity, some phytotoxins) or closure (abscisic acid, jasmonic acid, high ambient CO₂,

ozone, Pathogen Associated Molecular Patterns (PAMPs)) (Maurel *et al.*, 2015). Thus, stomatal aperture can adapt to daily fluctuations in evaporative demand or light, respond on the longer-term to soil water or atmospheric CO_2 availability and mediate elaborate plant strategies to counteract pathogen attacks.

Aquaporin expression in guard cells

As aquaporins can be found in essentially any plant tissue, it was no surprise to detect expression of aquaporins in guard cells of all higher plants examined, in species as diverse as Arabidopsis (Leonhardt et al., 2004; Prasch et al., 2015), sunflower (Helianthus annuus) (Sarda et al., 1997), broad bean (Vicia faba) (Sun et al., 2001), spinach (Spinacia oleracea) (Fraysse et al., 2005), maize (Zea mays) (Heinen et al., 2014), tree tobacco (Nicotiana glauca) (Smart et al., 2001) and Norway spruce (Picea abies) (Oliviusson et al., 2001). In contrast to certain plant organs such as seeds and pollen grains which express specific aquaporin isoforms, guard cells harbour PIPs and TIPs which are also expressed in other tissues (Smart et al., 2001; Leonhardt et al., 2004; Fraysse et al., 2005; Heinen et al., 2014; Prasch et al., 2015). Yet, many studies have shown that regulation of aquaporin expression in guard cells is linked to key components of stomatal regulation. In Arabidopsis guard cells, for instance, the transcript abundance of several PIPs was enhanced after 4 h in response to spray of ABA on the leaf surface (Leonhardt et al., 2004). In tree tobacco, expression of two TIP genes was reduced by drought after withholding water for 3-4 days (Smart et al., 2001). In sunflower, expression of a TIP showed diurnal variation, was transiently induced in response to water stress and in both cases was synchronized with stomatal closure (Sarda et al., 1997). In maize stomatal complexes, six out of seven PIP genes investigated showed a diurnal expression pattern with >3-fold higher expression in the morning than at night (Heinen et al., 2014). Finally, expression of two PIP and three TIP genes was reduced in an Arabidopsis mutant lacking a guard cell β-amylase and defective in starch degradation and stomatal opening (Prasch et al., 2015).

Cellular roles

Water transport

Initial evidence for a role of aquaporins in stomatal movements was rather indirect. For instance, the dose-dependent effects of extracellular calcium on stomatal aperture in broad bean leaf epidermal peals were tentatively associated to an activation and an inhibition of aquaporins by low and high calcium concentrations, respectively (Yang *et al.*, 2006). Shope & Mott (2006) also investigated broad bean guard cells and used the kinetics of osmotically induced changes in cell volume to estimate their apparent hydraulic conductivity. Interestingly, this parameter was sensitive to membrane trafficking inhibitors (cytochalasin D, wortmannin) provided that guard cells had received a hyperosmotic pretreatment. This was interpreted to mean that protein (aquaporin)-mediated transport of water

had been induced in these conditions. Consistent with this idea, the water permeability of isolated Arabidopsis guard cell protoplasts was enhanced twofold in response to a 10 µM ABA treatment (Grondin et al., 2015). The role of AtPIP2;1 aquaporin in this process was assessed using protoplasts from pip2;1 knock-out plants which showed a similar basal water permeability as wild types but lacked any response to ABA. In fact, AtPIP2;1 can be phosphorylated at a specific cytosolic site (Ser121) and activated by the Snf1-Related protein Kinase 2.6 (SnRK2.6, also named OST1) (Grondin et al., 2015). OST1 is itself released from inhibition by clade A Protein Phosphatases 2C (PP2C) such as ABA Insensitive 1 (ABI1), upon binding of ABA to RCAR/Pyr/PYL receptors which in turn capture PP2Cs (Cutler et al., 2010) (Fig. 1). The role of Ser121 phosphorylation in activation of AtPIP2;1 by ABA was definitely established by water transport measurements in guard cell protoplasts expressing phosphodeficient or phosphomimetic mutant forms of AtPIP2;1 at Ser121. Consistent with an ABA-dependent activation of AtPIP2;1, stomata

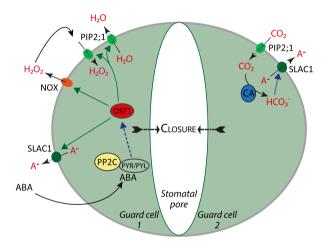


Figure 1. Simplified overview of implication of aquaporins in ABAand CO2-induced stomatal movements in Arabidopsis. The drawing represents two guard cells (light green) delineating a stomatal pore (white). Left (guard cell 1), under resting conditions, clade A Protein Phosphatase 2C (PP2C) family members, such as ABA Insensitive 1 (ABI1), act as negative regulators of ABA signalling, inhibiting the Snf1-Related protein Kinase 2.6 (SnRK2.6, also named OST1) via physical interaction, and leaving the S-type anion channel (SLAC1) with basal activity. Under drought, the ABA concentration in leaves increases and is perceived via Regulatory Component of ABA Receptor (RCAR) family members (Pyr/PYL). The ABA-induced formation of the RCAR/Pyr/PYL-PP2C complex breaks the PP2C-OST1 complex, thereby releasing active OST1 kinase. In turn, OST1 phosphorylates and activates several guard cell membrane proteins, such as the plasma membrane aquaporin AtPIP2;1 (PIP2;1), NADPH oxidases (NOX) such as RbohD, and the SLAC1 anion channel. Apoplastic H2O2 resulting from NOX activity may diffuse into the guard cell through AtPIP2;1 to trigger subsequent signalling events essential for stomatal closure. Release of anions through SLAC1 depolarizes the membrane and triggers cation efflux. The accompanying osmotic efflux of water mediated by AtPIP2;1 leads to a drop in guard cell turgor and to stomatal closure. **Right** (guard cell 2), AtPIP2;1 also physically interacts with plasma membrane β-Carbonic Anhydrase 4 (CA), facilitating the transmembrane diffusion of ambient CO2 into the guard cell. Resulting intracellular bicarbonate binds to and activates SLAC1, thereby promoting stomatal closure.

of *pip2;1* plants opened and closed normally in response to changing light or ambient CO₂ whereas they failed to close in response to ABA. Further, this response could be restored after expression of the phosphomimetic but not the phosphodeficient form of AtPIP2;1 at Ser121 (Grondin et al., 2015). We note that a role for PIPs in stomatal responses to ABA had previously been suggested by the finding that overexpression of a V. faba PIP1 in Arabidopsis leaves accelerated ABA-induced stomatal closure in a peeled epidermal assay (Cui et al., 2008). As indicated above, stomatal complexes of grasses have subsidiary cells which participate in stomatal movement via bulk water and ion flow into guard cells (Raschke & Fellows, 1971). It will be interesting to investigate whether aquaporins fulfil specific functions in these cells to assist stomatal movements. More generally, pharmacological and genetic experiments in algae (Chara corallina), moss (Physcomitrella patens) or higher plants (Raphanus sativus) have shown that aquaporins are not limiting for evaporation from outer cell wall surface (Tazawa & Okazaki, 1997; Lienard et al., 2008), as would occur in stomatal chambers.

H_2O_2 transport

Grondin et al. (2015) have suggested that the failure of pip2;1 guard cells to close in response to ABA may be because of combined defects in membrane water transport and hormonal signalling. In support for the latter hypothesis, pip2;1 guard cells lacked the typical intracellular accumulation of reactive oxygen species that occurs over the 30 min following exposure to ABA. H₂O₂ is produced in the apoplasm through ABAdependent activation of plasma membrane NADPH oxidases and plays a central role in ABA signalling (Murata et al., 2015). However, the sites of action of H_2O_2 and its modes of diffusion into the cells have remained undetermined. Aquaporins, which were shown to transport H2O2 after functional expression in yeast (Bienert et al., 2007; Dynowski et al., 2008), represent interesting candidates for this function (Fig. 1). This hypothesis is currently being investigated in our laboratory using the HyPer genetic probe as an intracellular reporter of H₂O₂ (Rodrigues et al., unpublished data).

CO₂ transport

Functional expression in yeast cells or *Xenopus* oocytes have indicated that some of the PIP isoforms expressed in maize or *Arabidopsis* guard cells (*Zm*PIP1;5, *Zm*PIP1;6, *At*PIP2;1) can transport CO₂ in addition to water (Heinen *et al.*, 2014; Wang *et al.*, 2016). The preferential localization of *Zm*PIP1;5 and *Zm*PIP1;6 at the plasma membrane (and not the chloroplast envelope) suggested a role in CO₂ signalling rather than CO₂ fixation (Fig. 1). This idea was recently corroborated by the finding that *At*PIP2;1 physically interacts with plasma membrane β-Carbonic Anhydrase 4 (βCA4) (Wang *et al.*, 2016). Co-expression in *Xenopus* oocytes of the two partners with Slow Anion Associated-Channel 1 (SLAC1) and activating protein kinases (OST1, CPK6 or CPK23) was necessary to confer on SLAC1 an enhancement by extracellular CO₂. This study led to a model whereby *At*PIP2;1 and βCA4 cooperatively facilitate the transmembrane diffusion of ambient CO_2 to enhance the intracellular concentration of bicarbonate (Fig. 1). Bicarbonate in turns binds to and activates SLAC1, thereby promoting stomatal closure. However, *pip2;1* guard cells showed a normal response to external CO_2 , likely because of a functional redundancy of *At*PIP2;1 with other PIP isoforms. Thus, this interesting model awaits confirmation in the plant.

Conclusion

Initial studies on the function of aquaporins in guard cells have focused on the key role of these proteins in membrane water transport. While this role definitely remains relevant, a few recent reports point to new aquaporin functions related to cell signalling, with potential significance beyond the context of stomatal regulation (Heinen et al., 2014; Grondin et al., 2015; Wang et al., 2016). A future challenge will be to determine how these functions lead to integration of aquaporins in the numerous signalling pathways acting in guard cells and how aquaporins may themselves create cross-talks between these pathways. Phosphorylation of AtPIP2;1 in guard cells is enhanced by ABA (Grondin et al., 2015) while gene expression of this and other PIPs can be regulated by H₂O₂ in leaves and roots (Hooijmaijers et al., 2012). Thus, it will be crucial to establish more precisely how the signalling pathways at work in guard cells can themselves act on the activity (or subcellular localization) of aquaporins. The functional redundancy of the numerous aquaporin isoforms expressed in guard cells may hinder the genetic validation of these analyses.

Whereas studies have so far focused on the role of PIPs, TIPs may also play key and original roles in stomatal responses. For instance, compartmental analyses of ion fluxes in osmotically challenged guard cells of *Commelina communis* have revealed that the tonoplast might be able to sense local osmotic gradients to promote vacuolar ion release. A tentative role in osmosensing has been assigned to aquaporins sitting in this membrane (MacRobbie, 2006).

WHOLE PLANT AQUAPORIN FUNCTIONS AND PLANT TRANSPIRATION

Principles of plant transpiration

Several steps possibly determine the transfer of water in the soil-plant-atmosphere continuum and, as a consequence, plant water use. For instance, reduced water equilibration in the rhizosphere can result in local dehydration and thereby a high soil hydraulic resistance (Draye *et al.*, 2010). The diffusion of water vapour from stomatal apertures is another strongly limiting process. It is directly determined by meteorological factors (e.g. vapour pressure deficit, leaf temperature) and stomatal aperture, which is itself governed by physiological (e.g. ABA) and environmental (e.g. light, ambient CO₂) variables. The previous section showed how aquaporins contribute to integrating some of these variables for optimized adjustment of stomatal aperture. By comparison to the rhizosphere and stomata, the hydraulic resistances of inner plant segments (roots, stems,

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leaves) are assumed to be much lower (Taiz & Zeiger, 1991). Whereas their direct impact on the rate of transpiration (*E*) is reduced, these hydraulic resistances crucially determine water potential gradients throughout the plant and, thereby, its water status. In the present section, we examine the mutual links existing between aquaporin activity in these organs and plant transpiration and/or stomatal conductance (g_s). Although not exhaustive, this section emphasizes some of the mechanisms at work for optimizing stomatal functioning and plant transpiration efficiency (i.e. the ratio of mass accumulation to transpiration).

Aquaporin genetic manipulation

The literature abounds with reports showing that genetic manipulation of aquaporins can dramatically alter E or g_s . Table 1 lists a representative subset of these reports. It shows that a decrease or enhancement of g_s by up to 30–40% could be observed in several studies. While most of these addressed the function of PIPs, it is of note that genetic manipulation of TIPs can also lead to alterations in E and g_s (Lin *et al.*, 2007; Sade *et al.*, 2009). In some cases, the effects of aquaporin genetic manipulation on root or leaf hydraulics were also characterized. Overall, the spectacular plant phenotypes support a crucial role of aquaporins in plant water relations. Yet, their interpretation with regard to stomatal regulation remains uncertain.

First, transgenic overexpression of an aquaporin gene or targeting of several aquaporin genes using antisense, RNAi or miRNA strategies can lead to a dramatic deregulation of the whole plant hydraulics which in turn may impact stomatal conductance. These difficulties can be circumvented using different approaches. For instance, grafting experiments were used by Sade et al. (2010) to uncouple aquaporin effects in roots and shoots. With respect to transgenic tomato plants that ectopically expressed tobacco NtAQP1, plants with wild-type root systems carrying transgenic shoots showed reduced E without significant difference in gs. To address guard cellspecific aquaporin functions, Sade et al. (2014a) have expressed NtAQP1 in transgenic Arabidopsis under the control of a KST1 stomata-specific promoter. Surprisingly, no alteration in gs was observed whereas expression of NtAQP1 using a photosynthetic tissue-promoter resulted in enhanced g_s . A limitation in these and other studies is because of heterologous aquaporin expression: the transgene-encoded aquaporin can escape tissue-specific molecular and cellular regulations targeting endogenous aquaporins and even lead to their deregulation (Jang et al., 2007).

Another difficulty may arise from confounding effects of aquaporin CO₂ transport activity. Table 1 shows that a covariation of g_s with mesophyll conductance to CO₂ (g_m) could be observed in all studies where the two parameters were investigated. The molecular and physiological mechanisms which possibly link these two parameters are as yet unknown (Flexas *et al.*, 2006; Flexas *et al.*, 2013). Nevertheless, it remains difficult to deduce the primary effects of the targeted aquaporin(s) based on such integrated phenotypes.

Overall, these genetic approaches provide significant and promising results to explore the integrated role of aquaporins

and their impact on transpiration. They may be refined, by using for instance tissue-specific complementation in corresponding aquaporin knock-out mutants. Such strategy was used in leaf veins (Prado *et al.*, 2013) but remains to be developed for stomatal expression of aquaporins. For now, a safe approach may be to relate whole plant and isolated stomata phenotypes. In these respects, the reduced transpiration rate of transgenic *Arabidopsis* expressing a *V. faba* PIP1 (*Vf*PIP1) could be connected to an enhanced stomatal closing response to ABA or darkness in a peeled epidermis assay (Cui *et al.*, 2008).

Root and shoot hydraulic conductance and transpiration

Consistent with the genetic studies discussed above, a large body of physiological data can explain how changes in hydraulics and related aquaporin activities occurring throughout the plant body can dramatically impact stomatal function. For instance, xylem vessel embolism leads to a drop in leaf water potential which mechanically induces stomatal closure. By facilitating embolism refilling, aquaporins may thus indirectly promote stomatal opening (Secchi & Zwieniecki, 2014). The living cells wrapping leaf veins, including xylem parenchyma cells and bundle sheath cells, also represent another hydraulic constriction for leaf water supply (Ache et al., 2010; Shatil-Cohen et al., 2011; Prado et al., 2013). As aquaporins critically determine the water permeability of bundle sheath cells, their down-regulation by ABA provides a powerful mechanism for promoting stomatal closure (Shatil-Cohen et al., 2011; Pantin et al., 2013). The dual effects of ABA on stomatal closure were elegantly demonstrated in Arabidopsis leaves by using an ost2 mutant which guard cells have become insensitive to ABA (Pantin et al., 2013). Yet, this mutant was able to close stomata in response to ABA through hydraulic regulation of leaf vein cells. Thus, in plants under water stress, ABA inhibits and enhances water transport (aquaporin activities), in bundle sheath and guard cells, respectively. These opposite effects both contribute to stomatal closure. Regulation of root hydraulic conductivity can also impact leaf water relations (Ehlert et al., 2009). In maize plantlets, pharmacological inhibition of root aquaporins revealed that the expansive growth of leaves was more sensitive than transpiration to root water transport. Yet, an inhibition of root hydraulic conductivity by 50% was able to reduce E by >50%, provided that the plant was under high evaporative demand (2.8 kPa VPD and 400 μ mol m⁻² s⁻¹ photosynthetic photon flux density) (Ehlert et al., 2009). Models that integrate effects of ABA on root and shoot hydraulics and stomatal aperture are needed to apprehend the full impact of the hormone on plant transpiration.

The interplay between transpiration and aquaporin regulation

Regulation of aquaporins by transpiration

In agreement with transgenic approaches in tomato (Sade et al., 2009), two recent studies on natural variation and water

Acuanorin	Origin plant species	Host plant species	Genetic alteration	Tissue specificity	Hydraulics and water relations	Photosynthesis (A_N)	ő	8 ^m	Reference
name					Percentage change	Percentage change with respect to wild type	'pe		
NtAQP1	Nicotiana tabacum	N. tabacum	Antisense	Whole Mole	$L_{\rm Dr}: -42 \ \Psi_{\rm stem}: -19 \ \Psi_{\rm leaf}: -10 \ E: -7$	n.d.	-32	n.d.	(Siefritz <i>et al.</i> ,
NtAQP1	N. tabacum	N. tabacum	Overexpr.	Whole	(Juguota) 2. – J2 (Juguota) n.d.	+136	+43	+64	(Uehlein
NtAQP1	N. tabacum	N. tabacum	Antisense	plant Whole	n.d.	-57	-33	-32	<i>et al.</i> , 2003) (Uehlein
H√PIP2;1	Hordeun vulgare	Oryza sativa	Overexpr.	plant Whole	n.d.	+18	+27	+40	<i>et al.</i> , 2003) (Hanba <i>et al.</i> ,
NtAQP1	N. tabacum	N. tabacum	Antisense	plant Whole	n.d.	-13	-30	-30	2004) (Flexas <i>et al.</i> ,
NtAQP1	N. tabacum	N. tabacum	Overexpr.	plant Whole	n.d.	+20	+40	+20	2006) (Flexas <i>et al.</i> ,
NtAOP1	N. tabacum	N. tabacum	RNAi	plant Whole	n.d.	- S	-15	-20	2006) (Uehlein
	- c		(Plant	- - -	, -	-	-	et al., 2008)
IJIJÍA	Vicia Jaba	Arabiaopsis thaliana	Uverexpr.	w nole plant	E: -Z1 (Drought)	n.a.	п.а.	n.a.	(Cui <i>et al.</i> , 2008)
NtAQP1	N. tabacum	N. tabacum	Overexpr.	Whole	Lp_r : +113 (Stress) Mesophyll P_c : +178 F : +25	+41	+39	*	(Sade <i>et al.</i> , 2010)
AtPIP1;2	A. thaliana	A. thaliana	T-DNA	Whole	n.d.	-23	Stomatal	-21	(Heckwolf
<i>Mc</i> MIPB	Mesembrvanthemum	N. tabacum	insertion Overexpr.	plant Whole	n.d.	+31	aperture : -25 +13	+35	<i>et al.</i> , 2011) (Kawase <i>et al.</i> ,
	crystallinum		-	plant	1	č	20	č	2013)
MAUTI	IV. IaDacarn	A. manana	Overexpr.	MESOPILYI	11.tl.	T2T	00+	+24	(Jaue et m., 2014a)
<i>At</i> PIP1;1 –5	A. thaliana	A. thaliana	miRNA	Whole	K_{leaf} : -50 K_{ros} : -32 Bundle sheath P_e : -67 Mesonhvill P_e : -68	-18	-21	-20	(Sade <i>et al.</i> , 2014b)
<i>At</i> PIP1;1-5	A. thaliana	A. thaliana	miRNA	Bundle sheath	$K_{\text{leaf}} : -62 \text{ Bundle sheath } P_{\text{f}} : -55$ Mesonhvll $P_{\text{f}} \cdot -80$	n.s.	n.s.	n.s.	(Sade <i>et al.</i> , 2014b)
PcPIP1;1-1;	Populus canescens	P. canescens	RNAi	Whole	$K_{\text{leaf:}} + 60 E$: +23	+37	+47	+133	(Bi <i>et al.</i> , 2015)
PgTIP1	Panax ginseng	A. thaliana	Overexpr.	Whole	E: +38	n.s.	+18	n.d.	(Lin et al.,
SITIP2;1	Solanum	S. lycopersicum	Overexpr.	plant Whole	E: n.s. (Control) E : +56–100 (Stress)	n.d.	n.d.	n.d.	2007) (Sade <i>et al.</i> ,

Table 1. Effects of aquaporin genetic manipulation of plant water relations and photosynthesis. The function of the indicated aquaporin gene was manipulated in the original or in a heterologous plant

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stress responses in grapevine (Vandeleur *et al.*, 2009; Pou *et al.*, 2013) have pointed to a positive relation between aquaporin expression and whole plant transpiration or g_s . Interestingly, this relation not only applies to PIPs which fulfil obvious roles in transcellular water transport and tissue hydraulics (Vandeleur *et al.*, 2009) but also to TIPs (Pou *et al.*, 2013). Because they rely on wild-type plants, these studies (Vandeleur *et al.*, 2009; Pou *et al.*, 2013) avoid the drawback of current aquaporin genetic manipulations. However, they could not elucidate the causal origin of the relationship between aquaporin expression and plant transpiration.

While the physiological studies discussed above establish a general frame for understanding the direct and indirect impacts of aquaporins on plant transpiration, a feedback effect of transpiration on aquaporin function is emerging. Establishing such effect is not trivial as transpiration has to be uncoupled from other types of parameters such as light or temperature that usually co-vary with transpiration, during diurnal cycles in particular. In relation to diurnal changes in root hydraulic conductivity of rice plants, OsPIP2;5 was shown to be specifically responsive to transpiration as its mRNA and protein accumulation in roots during the day could be reduced when shoots were exposed to high humidity (Sakurai-Ishikawa et al., 2011). A similar approach for uncoupling transpiration from light was developed in hybrid poplar (Laur & Hacke, 2013). In this case, an increase in PIP1 and PIP2 expression in roots could be induced by a low relative humidity treatment whereas g_s remained unchanged. Aquaporin expression in roots, root hydraulic conductivity and evaporative demand (based on meteorological factors) were also determined in rice plants grown under field conditions, and were nicely correlated, in good agreement with observations made in growth chamber experiments (Murai-Hatano et al., 2015). In these approaches, expression of five PIPs and a TIP showed a strong positive correlation to evaporation potential, whereas expression of a PIP and a TIP homolog, which seem to be associated with cell elongation, showed a negative correlation. The effects of a low relative humidity (i.e. high transpiration) are not restricted to root hydraulics. In Arabidopsis, a high evaporative demand resulted in an increase by > 3-fold in leaf hydraulic conductance (K_{leaf}) (Levin et al., 2007). In rice leaves, a coordinated up-regulation of several PIP and TIP genes could be observed as soon as 4 h after a dry air treatment (Kuwagata et al., 2012).

The signalling mechanisms which link plant transpiration to aquaporin activity in shoots and roots are as yet unclear. The rapid down-regulation of root hydraulics observed after shoot topping or defoliation may pertain to the shoot-to-root signalling involved (Liu *et al.*, 2014; Vandeleur *et al.*, 2014). This process was more specifically investigated in soybean and grapevine. It was proposed that a xylem-mediated hydraulic signal could be responsible for the change in root aquaporin expression observed within the 0.5–1 h following shoot topping (Vandeleur *et al.*, 2014). Conversely, the negative pressure (tension) present in xylem vessels of intact, transpiring plants could be perceived as an activating signal for aquaporin expression in root and shoot tissues.

Isohydric versus anisohydric plants

These recent studies may provide new ways to mechanistically understand the distinct behaviours of isohydric and anisohydric plants. The former ones have a conservative behaviour to maintain variations of leaf water potential at a minimum whereas the latter favour gas exchange at the expense of leaf water potential. Anisohydric plants have a more risky behaviour but are of potential agronomic interest as they can perform better than isohydric plants under mild water stress (Moshelion *et al.*, 2015). Reduced transpiration, which allows optimizing daily transpiration efficiency and long-term soil water availability, can also be an interesting agronomic trait. In a set of fieldgrown maize genotypes, reduced transpiration was associated to increased plant productivity in drought-prone environments (Messina *et al.*, 2015).

The current idea to explain plant anisohydric behaviour is that enhanced aquaporin expression and activity in roots and shoots promote whole plant hydraulic conductance thereby buffering water potential and favouring open stomata. Stomatal opening, as a consequence, promotes carbon fixation and plant growth (Sade et al., 2009; Vandeleur et al., 2009; Moshelion et al., 2015). For instance, an anisohydric grapevine cultivar (Chardonnay) was found, at variance with an isohydric cultivar (Grenache), to maintain the diurnal activation of root aquaporins under water stress to match the plant transpiration demand (Vandeleur et al., 2009). Thus, all mechanisms which provide a mutual coupling of tissue hydraulics and transpiration could be of prime importance for plant productivity. However, it is as yet unclear whether isohydric and anisohydric cultivars differ on direct aquaporin sensitivity to transpiration (see above) or on integrated ABA regulations. In particular, it is not yet known whether the guard cell mechanisms discussed in the first part of this review may differ between the two types of plants. Interestingly, some plant species such as grapevine or olive tree can switch between isohydric and anisohydric behaviours, depending on their environmental or developmental context (Moshelion et al., 2015). These materials will be useful to further explore the mechanistic bases of these two behaviours.

CONCLUSIONS

Although transpiration and aquaporins have long been identified as two key components influencing plant water status, it is only recently that their relations have been examined in detail. Multiple and unanticipated facets are currently emerging. In particular, recent studies indicate how guard cells can be used as a model to address new roles of aquaporins in cell signalling and movements. In addition, we now have a better understanding on how aquaporins throughout the plant exert multiple indirect effects on stomatal aperture. The latter impacts the leaf water status and carbon fixation, which in turn interfere with expansive growth and biomass accumulation. Future studies will have to further explore intimate links between aquaporins and plant growth (Maurel *et al.*, 2015). They may provide new directions for controlling or engineering the transpiration efficiency of crop plants, under replete or limiting water supply, thereby improving crop performance and productivity (Moshelion *et al.*, 2015).

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