

## REVIEW ARTICLE

## Function and regulation of phospholipid signalling in plants

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As an important metabolic pathway, phosphatidylinositol metabolism generates both constitutive and signalling molecules that are crucial for plant growth and development. Recent studies using genetic and molecular approaches reveal the important roles of phospholipid molecules and signalling in multiple processes of higher plants, including root growth, pollen and vascular development, hormone effects and cell responses to

environmental stimuli plants. The present review summarizes the current progress in our understanding of the functional mechanism of phospholipid signalling, with an emphasis on the regulation of  $\text{Ins}(1,4,5)P_3$ - $\text{Ca}^{2+}$  oscillation, the second messenger molecule phosphatidic acid and the cytoskeleton.

Key words: phospholipid, plant, signalling.

## BACKGROUND

Membranes act as barriers to hydrophilic molecules and ions because of the hydrophobic core of the phospholipid bilayer [1]. Phospholipid molecules are one of the main structural components of membranes, and they have emerged as important second messengers [2] to regulate plant growth and development and cellular responses to environmental change or stress [3].

Phosphorylations of the inositol ring of phosphatidylinositol are carried out by specific phosphoinositide kinases, including PI3Ks (phosphoinositide 3-kinases) [4], PI4Ks (phosphoinositide 4-kinases) [5] and PI5Ks (phosphoinositide 5-kinases) [6] at the D-3, D-4 or D-5 positions to generate  $\text{PtdIns}3P$ ,  $\text{PtdIns}4P$  or  $\text{PtdIns}5P$  respectively. Sequential phosphorylation by phosphoinositide 4-phosphate 5-kinase [7] or phosphoinositide 5-phosphate 4-kinase [8] will generate  $\text{PtdIns}(4,5)P_2$ , which is then hydrolysed by PLC (phospholipase C), resulting in the production of two important second messenger molecules:  $\text{Ins}(1,4,5)P_3$  and DAG (diacylglycerol) [2]. The inositol polyphosphates can be phosphorylated to generate  $\text{Ins}P_6$  or dephosphorylated by inositol polyphosphate phosphatases, which are classified into four groups on the basis of the position of the phosphates [9]. In addition, PLD (phospholipase D) hydrolyses phospholipids at the terminal phosphodiester bond and generates PA (phosphatidic acid) [10] (Figure 1).

The presence of many isoforms of the key enzymes in phosphoinositide metabolism has been demonstrated. Some 12 PI3/4K-domain-containing proteins are predicted to be PI4K genes [11], and there are 15 isoforms of PIP5Ks (phosphoinositide phosphate 5-kinases) [11], 15 isoforms of 5PTases (inositol polyphosphate 5-phosphatases) [12], 12 PLD isoforms [13], 12 PI4K isoforms [10] and nine PLC isoforms [10,14] in *Arabidopsis*. In rice, a monocotyledon, the distribution and functions of relevant isoforms have been less studied. An analysis showed that there are 17 PLD members [15] and four PLC members [14] in rice.

Our preliminary analysis using BLAST and CLUSTW shows that there are ten PIP5Ks, 11 PI4Ks and 20 5PTase members in rice (Figure 1).

In the present review, we focus on the functions and mechanisms of PIP5K, PLC, 5PTase, PLD and the phospholipid molecules  $\text{PtdIns}4P$ ,  $\text{PtdIns}(4,5)P_2$ ,  $\text{Ins}(1,4,5)P_3$  and PA, as well as  $\text{Ca}^{2+}$ , in plant growth and development.

## CRITICAL EFFECTS OF PHOSPHOLIPIDS IN PLANT GROWTH AND STRESS RESPONSES

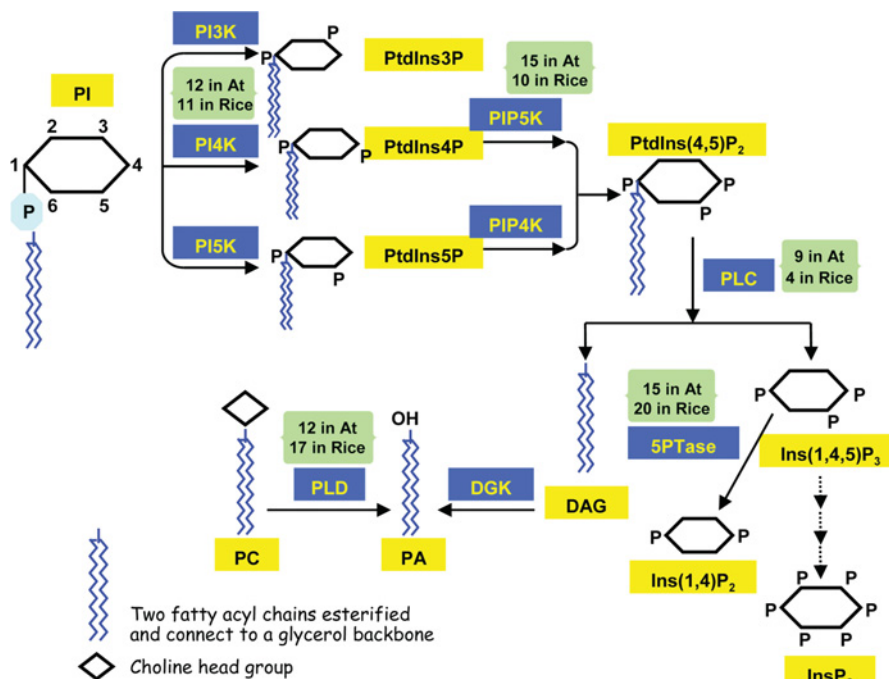
## Root growth and root hair patterning

Mutation in *XIPOTLI*, which encodes a PEAMT (S-adenosyl-L-methionine:phosphoethanolamine N-methyltransferase) and is critical in phosphatidylcholine biosynthesis, resulted in a short primary root and short epidermal cells due to the cell death induced by decreased phosphatidylcholine content [16]. *Arabidopsis* lines with a deficiency of *INT1* (*INOSITOL TRANSPORTER 1*) show increased intracellular *myo*-inositol concentrations and reduced root growth, which is possibly due to sequestration of inositol in vacuoles [17]. *Arabidopsis* PIP5K9 negatively regulates sugar-mediated root cell elongation through interaction with a cytosolic invertase, CINV1 [18]. Conversely, suppressed *AtIPK2α* (*Arabidopsis thaliana* inositol polyphosphate kinase 2α) expression and application of exogenous  $\text{Ins}(1,4,5)P_3$  lead to enhanced root growth [19].

Seedlings treated with the PLD inhibitor butan-1-ol or deficient in *PLDζ2* display suppressed primary root elongation and inhibited lateral root formation [20], and PLD-derived PA production is an early signalling event during adventitious root formation induced by auxin and NO in cucumber explants [21]. In addition, accumulated  $\text{PtdIns}(4,5)P_2$  and  $\text{Ins}(1,4,5)P_3$  under *SAC9* deficiency result in shorter primary roots and fewer lateral roots

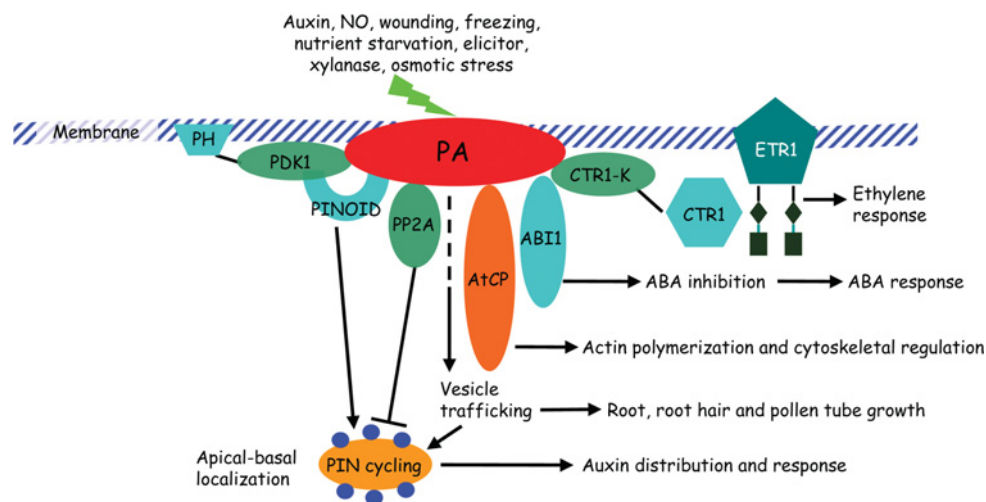
Abbreviations used: ABA, abscisic acid; AtCP, actin-capping protein; AtIPK2, *Arabidopsis thaliana* inositol polyphosphate kinase 2; AVR4, avirulence 4; CTR1, constitutive triple response 1; DAG, diacylglycerol; DGK, DAG kinase; F-actin, filamentous actin; IAA, indoleacetic acid;  $\text{IP}_3R$ ,  $\text{Ins}(1,4,5)P_3$  receptor; LOX2, lipoxygenase 2; LPP, lipid phosphate phosphatase; MAPK, mitogen-activated protein kinase; OsPIP, *Oryza sativa* phosphoinositide phosphate kinase; PA, phosphatidic acid; PDK1, phosphoinositide-dependent kinase 1; PI3K, phosphoinositide 3-kinase; PI4K, phosphoinositide 4-kinase; PIPK, phosphoinositide phosphate kinase; PIP5K, phosphoinositide phosphate 5-kinase; PLC, phospholipase C; PLD, phospholipase D; PP2A, protein phosphatase 2A; 5PTase, inositol polyphosphate 5-phosphatase; RACK1, receptor for activated C-kinase 1; SNX1, sorting nexin 1; ZmPLC1, *Zea mays* PLC1.

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**Figure 1** The phosphoinositide metabolic pathway in higher plants

The isoform numbers of the key enzymes in *Arabidopsis* (At) and rice are indicated. Abbreviations: PI, phosphoinositide; PI5K, phosphoinositide 5-kinase.



**Figure 2** PA functions in hormone responses and growth of roots, root hairs and pollen tubes

PA accumulates in response to treatment with auxin and NO, wounding, freezing, nutrient starvation, elicitor, xylanase, and osmotic stress. PA could bind CTR1 and block the interaction with ethylene receptor ETR1 to regulate the ethylene response. PA binds and decreases the activity of ABI1, resulting in the insensitive response to ABA treatment. PA promotes actin polymerization by binding to and inhibiting the activity of AtCP. PA regulates cycling of PIN proteins through directly mediating vesicle trafficking, or binding and mediating the activity of auxin transport upstream proteins PDK1, PINOID and PP2A. Abbreviations: CTR1-K, CTR1 kinase domain; PH, pleckstrin homology domain.

[22]. PA takes part in root growth, root hair and pollen tube patterning by regulating cytoskeleton and vesicle trafficking [20,21,23] (Figure 2).

Regarding the root hair pattern, a deficiency of *AtSfh1*, which encodes a PITP (phosphatidylinositol transfer protein), compromises polarized root hair expansion in a manner that coincides with a loss of tip-directed PtdIns(4,5)P<sub>2</sub> [24]. Decreased AtIPK2 $\alpha$  and PLD $\zeta$ 1 expression confers insensitivity of root hair growth to calcium depletion [19] and leads to random initiation of root

hairs [25] respectively, whereas increased *PLD $\zeta$ 1* expression induces branched and swollen root hairs [25]. Treatment with a PI3K-specific inhibitor inhibits root hair growth by regulating PtdIns3P-mediated vesicle trafficking and ROS (reactive oxygen species) production [26]. The double mutant *PI4K $\beta$ 1/PI4K $\beta$ 2* displays aberrant root hair morphologies due to altered calcium signalling [27], whereas mutation in a phosphoinositide 4-phosphate phosphatase gene (*rhd4*) results in shorter and morphologically aberrant root hairs due to the

altered accumulation of PtdIns4P together with PI4K $\beta$ 1 in membrane compartments [28]. Studies by two independent groups have shown that *Arabidopsis* PIP5K3 is essential for root hair formation through its production of PtdIns(4,5) $P_2$  at the root hair apex. Furthermore, reduced expression of PIP5K3 caused significantly shortened root hairs, whereas overexpression resulted in increasingly deformed root hairs [29,30]. In *Medicago truncatula*, inhibition of PI3K and PI-PLC (phosphatidylinositol-specific PLC) prevented root hair curling and the formation of infection threads [31].

### Pollen development

Phospholipid signalling is involved in pollen development, including pollen viability, germination and pollen tube elongation, through regulation of the production of PtdIns3P, PtdIns4P, PtdIns(3,5) $P_2$ , PtdIns(4,5) $P_2$ , PA, Ins(1,4,5) $P_3$  and downstream  $[Ca^{2+}]_{cyt}$  levels. Studies reveal that PI3K and PI4K are involved in regulation of pollen viability through different mechanisms. PI3K regulates vacuole reorganization and nuclear division in pollen grains, possibly by regulating the products PtdIns3P and PtdIns(3,5) $P_2$ , which are essential for protein targeting to vacuoles and normal vacuole function and morphology respectively [32], whereas PI4K $\gamma$ 1 performs crucial intracellular trafficking regulation for tapetum and microspore development [33]. In addition, PtdIns(4,5) $P_2$  leads to increased  $[Ca^{2+}]_{cyt}$ , transient growth perturbation and inhibition of apical secretion, whereas Ins(1,4,5) $P_3$  treatment causes a transient  $[Ca^{2+}]_{cyt}$  increase of similar magnitude and stimulates apical secretion and severe growth perturbation, indicating the different targets of these two molecules [23]. Recent reports have shown that deficiency of two *Arabidopsis* pollen-expressed PIP5Ks, PIP5K4 and PIP5K5, reduced pollen germination and pollen tube elongation, which is consistent with the effect of their product, PtdIns(4,5) $P_2$  [34,35]. However, overexpression of either PIP5K4 or PIP5K5 results in multiple pollen tip branching, which is thought to occur by regulating membrane trafficking and apical pectin secretion [34,35].

PA plays an important role in pollen tube polarity by regulating microfilament polymerization and membrane transport. A reduced level of PA leads to inhibited apical plasma membrane recycling and formation of thick, but non-directional, microfilaments [23]. PLD-derived PA stimulates pollen germination and tube elongation [36] and expands the pollen tube apical region [37] (Figure 2).

In addition, because both PtdIns(4,5) $P_2$  and DAG are accumulated in an apical domain of the plasma membrane at the pollen tube tips, disruption of this domain causes defective pollen tube growth, which highlights the importance of maintaining an apical domain enriched in PtdIns(4,5) $P_2$  and DAG for polar cell growth [38].

### Vascular development

Although several studies have demonstrated the role of phospholipid signalling in vascular growth and development, the mechanism is still unclear. CVP2, encoding a type I inositol polyphosphate 5-phosphatase 6 (*At5PTase6*), promotes vascular cell proliferation and prevents the termination of premature veins (the *cvp2*-knockout mutant shows an open cotyledon vein [39]). The type II *At5PTase13* regulates cotyledon vein formation by regulating auxin homeostasis [40]. FRA3, a type II 5PTase gene, plays an essential role in secondary wall synthesis in fibre cells and xylem vessels, and the *fra3*-knockout mutant contains higher amounts of PtdIns(4,5) $P_2$  and Ins(1,4,5) $P_3$  and shows reduced thickness of the secondary wall and decreased stem strength

owing to the disruption in actin organization [41]. AtSAC1 exhibits phosphatase activity toward PtdIns(3,5) $P_2$ , and mutation of *AtSAC1* causes a dramatic decrease in the cell wall thickness of fibre cells and vessel elements, resulting in a weak stem [42], suggesting the important effect of the PtdIns(3,5) $P_2$  pool in the formation of aberrant cell shape.

### Hormone effects

Altered phospholipid signalling shows changed responses to various hormones. Seedlings deficient in *5TPase1*, *5PTase2* [43] and *cvp2* [39] are hypersensitive to exogenous ABA (abscisic acid), whereas a deficiency in *5PTase13* causes insensitivity to ABA in seed germination [40]. PA, generated by PLD $\alpha$ 1, binds to ABI1 and decreases the phosphatase activity of ABI1, resulting in the insensitivity of stomatal closure of PLD $\alpha$ 1-null mutant leaves to ABA treatment [44] (Figure 2). In addition, PLD $\alpha$ 1 directly interacts with and inactivates the  $G_o$  subunit of heterotrimeric G-proteins to mediate ABA inhibition of stomatal opening [45], and suppression of rice *PLD $\beta$ 1* results in reduced sensitivity to ABA in seed germination [15].

Ins(1,4,5) $P_3$  oscillation mediates auxin transport and is involved in gravistimulation responses [46]. *Arabidopsis* *5PTase13* and *PLD $\zeta$ 2* regulate the cotyledon vein [40] and root development [20] respectively by modulating auxin synthesis and transport. *PLA* activity is rapidly induced by auxin within 2 min [47], and *Arabidopsis* *sPLA2* regulates shoot gravitropism by regulating auxin-induced cell elongation [48].

Besides auxin and ABA, involvement of phospholipid signalling in other hormone cascades is seldom reported. Testerink et al. [49] demonstrated that PA regulates CTR1 activity to influence ethylene signalling. Lin et al. [50] pointed out that phospholipid signalling could be regulated by multiple hormones, hinting at the cross-talk between them.

### Stress responses

The involvement of phospholipid signalling in the cell responses to stress stimuli, including salt, osmotic, temperature and pathogen stressors, has been well demonstrated, and multiple members of the phosphoinositide pathway are supposed to mediate the stress responses through different mechanisms.

#### Salt stress

Salinity is a major stress that threatens plant growth and crop production. *Arabidopsis* plants show rapidly increased PtdIns(4,5) $P_2$  synthesis in response to treatments with NaCl, KCl and sorbitol [51], suggesting an important role of PtdIns(4,5) $P_2$  in plant salt tolerance. PtdIns(4,5) $P_2$  and its derivative Ins(1,4,5) $P_3$  are accumulated in plants under salt, cold and osmotic stresses, and *sac* (suppressor of actin) mutants harbour accumulated PtdIns(4,5) $P_2$  and Ins(1,4,5) $P_3$  and show a constitutive stress response [22]. Consistent with this, pharmacological studies using U73122 (a specific inhibitor of PLC) have shown that, in salt-treated seedlings, PLC activity, as well as Ins(1,4,5) $P_3$  and calcium, are necessary for the expression of pyrroline-5-carboxylate synthetase and resultant pyrroline levels [52].

Recent studies revealed that salt-stress-induced association of PtdIns(4,5) $P_2$  with clathrin-coated vesicles [53] and PI3K-mediated endocytosis are necessary for plant salt tolerance [54], highlighting the importance of vesicle trafficking in the salt response. Multiple PLDs, including PLD $\alpha$ 1,  $\alpha$ 3,  $\delta$  and  $\epsilon$ , are required for salt and water deficit tolerance in *Arabidopsis*, possibly by regulating PA production and membrane lipid compositions [55,56].

### Drought stress

Drought directly leads to water deficiency and inhibits plant growth. Studies have shown that members of both PLC and PLD families are involved in drought tolerance. Expression of maize *ZmPLC1* (*Zea mays* PLC1) is up-regulated under dehydration and this enhanced expression improves the drought tolerance of maize [57]. Expression pattern analysis revealed that six of nine *Arabidopsis* PLC genes are stimulated by dehydration [14], suggesting a similar function of *Arabidopsis* PLCs to that of *ZmPLC1*. PLDs and the derived PA are also widely involved in plants' drought tolerance. Both *PLDα1* and *PLDα3* are necessary for drought tolerance in *Arabidopsis* [58]; however, the regulatory mechanisms are different. *PLDα1* mediates early drought response and ABA signalling [59], whereas *PLDα3* is involved in promoting root growth and stress avoidance under hyperosmotic conditions [55]. *PLDδ* is accumulated under dehydration and contributes to the dehydration-induced PA accumulation [60], whose knockout mutants display hypersensitivity to water deficit [56]. Recently, Perera et al. [61] showed that *Arabidopsis* transgenic plants overexpressing mammalian type I 5PTase present significantly enhanced drought tolerance.

### Cold stress

A transcriptomic analysis of cold-treated *Arabidopsis* suspension cells in the presence of U73122 or ethanol, the inhibitors for PLC and PLD respectively, revealed that both PLC- and PLD-mediated signal pathways are activated simultaneously, which leads to the activation of different clusters of cold-response genes [62]. Genetic analysis reveals that different PLD isoforms have opposite effects in cold response. *Arabidopsis* plants that are deficient for *PLDα1* show improved tolerance to freezing through activating the cold-responsive genes and increasing osmolyte accumulation [63,64], whereas deficiency of the membrane-associated *PLDδ* rendered hypersensitivity to freezing (*PLDδ* overexpression resulted in increased freezing tolerance [65]). Suppression of *PLDα1* decreases phospholipid hydrolysis and PA production in both freezing and post-freezing phases, whereas ablation of *PLDδ* increases lipid hydrolysis and PA production in post-freezing recovery [66]. Chilling and freezing result in the increased  $\text{Ins}(1,4,5)\text{P}_3\text{-Ca}^{2+}$  influx [67] and transgenic plants overexpressing mammalian 5PTase display a ~30% decrease in the rapid transient  $\text{Ca}^{2+}$  peak in response to cold or salt stimuli [61].

### Oxidative stress

$\text{H}_2\text{O}_2$  treatment activates the PLD activity, and deficiency of *PLDδ* displays hypersensitivity to  $\text{H}_2\text{O}_2$ -induced cell death, suggesting a crucial role of PA in  $\text{H}_2\text{O}_2$ -induced cell death [68]. This is consistent with the fact that PA could trigger an oxidative burst through activating a MAPK (mitogen-activated protein kinase) cascade [69] and indicates PA as a possible important regulator of Rop-regulated ROS generation to mediate the process of cell death [70]. In addition, PI3K can stimulate plasma membrane endocytosis and produce ROS, subsequently inducing root hair curling [31].

### Other stress

Wounding triggers a rapid activation of PLD-mediated phospholipid hydrolysis. The  $\text{Ca}^{2+}$  increase upon wounding mediates the translocation of PLD protein from cytosol to membrane [71]. Multiple members of PLD are activated by wounding, and

suppression of *PLDα* by an antisense approach decreases the wound-induced expression of *LOX2* (lipoxygenase 2), suggesting that *LOX2* is probably a downstream target through PLD-mediated production of jasmonic acid [72]. MAPK signalling is activated by wounding in soya bean seedlings, and wound-induced activation of MAPK signalling is suppressed when PA production is inhibited with n-butanol, which indicates that PA acts as a second messenger in wound-induced MAPK signalling [73]. In addition, *PLDζ2* plays a key role in phosphate cycling, and *pldζ2* mutants exhibit hypersensitivity to phosphate starvation [74,75]. *PLDε* regulates the root surface area to improve the nitrogen uptake and utilization [76].

Besides abiotic stress, phospholipid signalling is also regulated by biotic stress. The *AVR4* (avirulence 4) gene encodes an elicitor protein and displays resistance in tomato plants carrying the *Cf-4* resistance gene to the fungus *Cladosporium fulvum*. In response to the infection, PA is rapidly produced in a few minutes when *AVR4* interacts with *Cf-4*, indicating that the PA accumulation is an early response in the *Cf-4*–*AVR4* interaction [77].

### Others

In addition to the above-mentioned developmental processes and stress responses, phospholipid signalling is involved in embryo development, stomata closure, and light and sugar signal transduction. Deficiency of PECT (phosphorylethanolamine cytidyltransferase), a rate-limiting enzyme in phosphatidylethanolamine biosynthesis, results in embryo abortion before the octant stage, delays embryo maturation and reduces seed fertility [78]. Deficiency of LPAT (lysophosphatidyl acyltransferase), which is responsible for PA biosynthesis, caused embryo lethality (arrested embryo at the globular stage [79]).

The *Arabidopsis* 5PTase13 deficiency mutant (*5pt13*) displays shorter hypocotyls under blue light [80]. Mutants *5pt1* and *5pt2* grow faster and have elongated hypocotyls in the dark [43]. The *ITPK-1* (inositol 1,3,4-trisphosphate 5/6-kinase) knockout mutant is hypersensitive to red light, similar to the *csn* (COP9 signalosome) mutant [81]. In addition,  $\text{PtdIns}(4,5)\text{P}_2$  and  $\text{Ins}(1,4,5)\text{P}_3$ -mediated  $\text{Ca}^{2+}$  oscillation are important for stomatal opening [82,83]. *Arabidopsis* 5PTase13 regulates SnRK1 (sucrose non-fermenting-1-related kinase) activity, which differs according to nutrient availability, and deficiency of *5PTase13* results in a shortened root under no-nutrient or low-nutrient conditions and is less sensitive to sugar and insensitive to ABA [84].

The altered growth patterns of the loss-of-function knockout mutant and transgenic lines with overexpression or decreased expression of phospholipid signalling-related genes are summarized in Table 1.

## REGULATION OF PHOSPHOLIPID SIGNALLING

### Regulation of key enzymes and molecules in phosphoinositide signalling by hormones and environmental stimuli

Studies of expression patterns of genes encoding key enzymes in phosphoinositide signalling employing DNA chip technology and RT (reverse transcription)–PCR reveal that the expression of multiple enzymes is widely and differentially regulated by various hormones and abiotic stressors, especially in the families of PIPK (phosphoinositide phosphate kinase), PLD and 5PTase [50]. Of nine *Arabidopsis* PLC genes, four are induced by salt, three are induced by ABA, four respond to cold, and six are stimulated by dehydration [14]. *Arabidopsis* PLD is activated by various hyperosmotic stresses, including high salinity [93], dehydration [60] and freezing [63,65], as

**Table 1** Altered growth of loss-of-function knockout mutants (KO) and transgenic lines with overexpression (OE) or decreased expression (DE) of phospholipid signalling-related genesUnless indicated otherwise, results are from *Arabidopsis*.

Gene	Protein	Type	Phenotype	Reference(s)
<i>VSP34</i>	PI3K	KO	Short root hairs, and decreased pollen viability, germination and pollen tube growth	[26,32]
<i>PI4Kβ1/PI4Kβ2</i>	PI4Kβ1, PI4Kβ2	KO	(Double mutant) Aberrant root hair morphology	[27]
<i>PI4Kγ1</i>	PI4Kγ1	KO	Decreased pollen viability, abnormal tapetum and microspore development	[33]
<i>RHD4</i>	PI4P phosphatase	KO	Short and morphologically aberrant root hairs	[28]
<i>PIP5K3</i>	PIP5K3	DE	Shorter root hair	[29,30]
		OE	Deformed root hair	
<i>PIP5K4</i>	PIP5K4	KO	Reduced stomatal opening, and impaired pollen germination, tube growth and polarity	[34,35,82]
		OE	Perturbed pollen tube growth, multiple pollen tip branching	
<i>PIP5K5</i>	PIP5K5	OE	Multiple pollen tip branching	[34]
<i>PIP5K9</i>	PIP5K9	OE	Shortened primary root	[18]
<i>SAC1 (fra7)</i>	SAC domain phosphoinositide phosphatase 1	KO	Weak stem, decreased cell wall thickness	[42]
<i>SAC9</i>	SAC domain phosphoinositide phosphatase 9	KO	Shorter primary root and fewer lateral roots	[22]
<i>DAD1</i>	PLA1	KO	Defects in anther dehiscence, pollen maturation and flower opening	[85]
<i>AtsPLA2β</i>	AtsPLA2β	DE	Shortened leaf petioles and stems, delayed light-induced stomatal opening	[48,86]
		OE	Prolonged leaf petioles and inflorescence stems, faster stomatal opening	
<i>AtPLC1</i>	AtPLC1	DE	Insensitive to ABA in seed germination and growth	[87]
<i>ZmPLC1</i>	ZmPLC1	OE	Improved drought tolerance	[57]
<i>NtPLC3</i>	NtPLC3	OE	Inhibited pollen tube growth	[38]
<i>OsPLDβ1</i>	OsPLDβ1	KO	Reduced sensitivity to ABA during seed germination	[15]
<i>PLDα1</i>	PLDα1	KO	Decreased wound-induced synthesis of jasmonic acid, decreased drought tolerance, enhanced seed quality after storage, insensitive response of stomatal closure to ABA, enhanced sensitivity to high salinity, increased freezing tolerance	[56,58,64,65,72,88]
		OE	Increased sensitivities to salinity and water deficiency, later flowering in drought conditions	[55]
<i>PLDε</i>	PLDε	KO	Decreased sensitivities to salinity and water deficiency, earlier flowering in drought conditions	[76]
		OE	Decreased root growth and biomass accumulation, decreased lateral root elongation	
<i>PLDδ</i>	PLDδ	KO	Increased root growth and biomass accumulation, increased lateral root and root hair elongation	[68,89]
		OE	Increased sensitivity to H <sub>2</sub> O <sub>2</sub> -induced cell death, sensitive to freezing	
		OE	Increased freezing tolerance	
<i>PLDζ1</i>	PLDζ1	OE	Branched and swollen root hairs	[25]
		DE	Random initiation of root hairs	
<i>PLDζ1/PLDζ2</i>	PLDζ1, PLDζ2	KO	(Double mutant) Hypersensitive to phosphate deficiency in root growth	[74,75]
<i>PLDζ2</i>	PLDζ2	KO	Suppressed primary root elongation and inhibited lateral root formation, less sensitive to auxin, reduced root gravitropism	[20]
		OE	Enhanced primary root growth, root gravitropism, hypersensitive to auxin.	
<i>5PTase1/5PTase2</i>	5PTase1, 5PTase2	KO	(Double mutant) Faster germination and longer hypocotyl in the dark, hypersensitive to ABA	[43]
<i>5PTase5</i>	5PTase5	KO	Disrupted root-hair tip growth	[90]
<i>CVP2</i>	5PTase6	KO	Open reticulum and increased free vein endings	[39]
<i>5PTase11</i>	5PTase11	KO	Slower germination and decreased hypocotyl growth when grown in the dark	[91]
<i>FRA3</i>	5PTase12	KO	Dramatic reduction in secondary wall thickness and a concomitant decrease in stem strength	[41]
<i>5PTase13</i>	5PTase13	KO	Defect in development of the cotyledon vein, shortened hypocotyls and expanded cotyledons under blue light, hypersensitive to sugar and ABA in seed germination	[40,80,84]
<i>Itpk-1</i>	ITPK-1	KO	Decreased hypocotyl length under red light	[81]
<i>OsITL1</i>	OsITPK1	OE	(In tobacco) Decreased tolerance to NaCl during germination and seedling development.	[92]
<i>ATS2</i>	LPAAT (lysophosphatidic acid acyltransferase β)	KO	Embryo lethality	[79]
<i>XIPOTL1</i>	PEAMT	KO	Short primary root and induced cell death	[16]
<i>PECT</i>	PECT	KO	Embryo abortion before the octant stage, delayed embryo maturation and reduced seed fertility	[78]
<i>INT1</i>	INT1	KO	Reduced root length	[17]
<i>AtSth1</i>	PITP	KO	Short root hairs	[24]

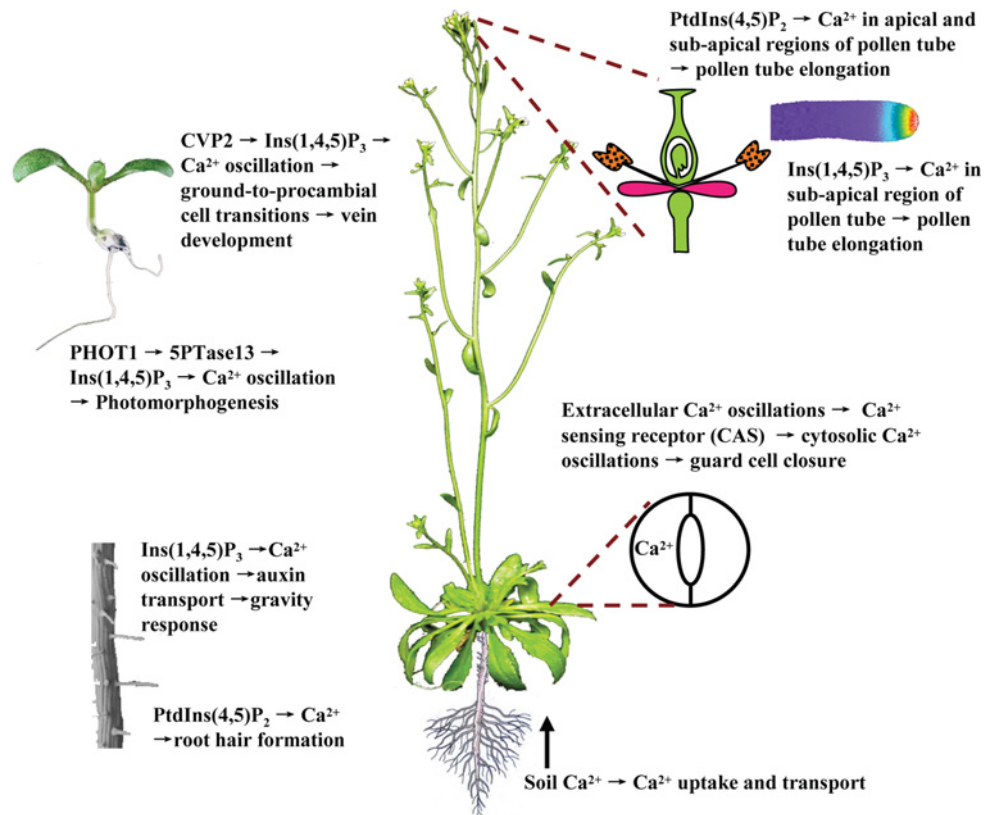
well as ABA [44]. Similarly, many rice *PLD* genes are induced by salt and drought stress [15]. *PIP5K* family genes are reported to be stimulated by water stress and ABA [50,59,94] and repressed by cold [18]. In addition, the plant hormone auxin is broadly related to phosphoinositide signalling, inducing the expression of *AtIPK2β* [95] and *PLDζ2* [20], and suppressing the expression of *5PTase13* [40].

Some phospholipid molecules, such as PA, PtdInsP and PtdIns(4,5)P<sub>2</sub>, are accumulated after treatment with auxin, NO [21], wounding [72,96], freezing [63], nutrient starvation [74,97], elicitor [98], xylanase [99] or under osmotic stress [100].

### Ins(1,4,5)P<sub>3</sub>-mediated Ca<sup>2+</sup> oscillation

PtdIns(4,5)P<sub>2</sub> throughout the membrane establishes the basis for rapid and transient increases of Ins(1,4,5)P<sub>3</sub> in response to environmental stimuli, which leads to the rapid oscillation of Ca<sup>2+</sup>. Many studies have shown that Ins(1,4,5)P<sub>3</sub>-mediated Ca<sup>2+</sup> influx is important for plant cell growth (Figure 3). In *cvp2* mutants, the Ins(1,4,5)P<sub>3</sub> levels increase approx. 3-fold compared with wild-type, and Ins(1,4,5)P<sub>3</sub> releases Ca<sup>2+</sup> from internal stores to tightly control the number of ground-to-procambial cell transitions and to regulate the vein formation [39]. The *5pt13* mutant has a





**Figure 3** The  $\text{Ca}^{2+}$  oscillation, mediated by  $\text{PtdIns}(4,5)\text{P}_2$  and  $\text{Ins}(1,4,5)\text{P}_3$ , is crucial for multiple processes of plant growth including root tropism, root hair formation, cotyledon vein development, photomorphogenesis, guard cell closure, pollen tube elongation and hormone effects

higher  $\text{Ins}(1,4,5)\text{P}_3$  level compared with wild-type and mediates the effects of blue light through  $\text{Ins}(1,4,5)\text{P}_3$ -mediated  $\text{Ca}^{2+}$  oscillations [80].

Gravistimulation induced differential lateral IAA (indoleacetic acid) transport, which is consistent with the changed  $\text{Ins}(1,4,5)\text{P}_3$  levels in maize and oat pulvini [46]. Previous studies have shown that the level of  $\text{Ins}(1,4,5)\text{P}_3$  increases approx. 3-fold in the first 5 min of gravistimulation, and the second peak of  $\text{Ins}(1,4,5)\text{P}_3$  appears by 15–20 min [101]. After 30 min, the  $\text{Ins}(1,4,5)\text{P}_3$  level returns to baseline [101], indicating that the oscillation of  $\text{Ins}(1,4,5)\text{P}_3$  is a key component of the response to gravity stimuli.

Environmental stimuli also evoke a transient increase of  $\text{Ins}(1,4,5)\text{P}_3$  levels, leading to the following question: is the long-term increase or decrease of the  $\text{Ins}(1,4,5)\text{P}_3$  levels associated with cell growth? The shorter hypocotyls of the *5pt13* mutant under blue light provide evidence that a higher  $\text{Ins}(1,4,5)\text{P}_3$  level, and hence increased  $[\text{Ca}^{2+}]_{\text{cyt}}$  under blue light irradiation, regulates cell growth [80]. Similarly, in maize pulvini, the inositol lipids' rate of turnover is associated with an increase in membrane biogenesis and  $\text{Ins}(1,4,5)\text{P}_3$ - $\text{Ca}^{2+}$  release from internal stores to initiate cell growth upon gravistimulation [102].

The  $\text{PtdIns}(4,5)\text{P}_2$ - $\text{Ca}^{2+}$  gradient plays a key role in pollen tube growth [103] and root hair formation [24]. Because  $\text{Ins}(1,4,5)\text{P}_3$ - $\text{Ca}^{2+}$  influx across the plasma membrane is required to maintain the  $\text{Ca}^{2+}$  gradient in the pollen tube cell and  $\text{Ins}(1,4,5)\text{P}_3$ -sensitive  $\text{Ca}^{2+}$  channels have not been identified [10], it was speculated that the function of  $\text{Ca}^{2+}$  influx was a direct result of  $\text{Ins}(1,4,5)\text{P}_3$  in pollen tube growth [38]. When  $\text{PtdIns}(4,5)\text{P}_2$  and  $\text{Ins}(1,4,5)\text{P}_3$  were loaded into an *Agapanthus* pollen tube,  $\text{Ins}(1,4,5)\text{P}_3$  was more effective than  $\text{PtdIns}(4,5)\text{P}_2$  [23], indicating that  $\text{Ins}(1,4,5)\text{P}_3$ - $\text{Ca}^{2+}$  and  $\text{PtdIns}(4,5)\text{P}_2$ - $\text{Ca}^{2+}$  are

spatially different.  $\text{PtdIns}(4,5)\text{P}_2$ - $\text{Ca}^{2+}$  is increased in both apical and sub-apical regions, whereas  $\text{Ins}(1,4,5)\text{P}_3$ - $\text{Ca}^{2+}$  is increased mainly in sub-apical regions [23], indicating that both  $\text{PtdIns}(4,5)\text{P}_2$  and  $\text{Ins}(1,4,5)\text{P}_3$  cause transient increases in  $\text{Ca}^{2+}$  levels; however, the effects are markedly different.

Transient and rapid increase in intracellular  $\text{Ca}^{2+}$  have been demonstrated in response to abiotic stresses, including cold, salt and osmotic stress [104,105], and  $\text{Ins}(1,4,5)\text{P}_3$ -mediated  $\text{Ca}^{2+}$  oscillation is also involved in the response to these stimuli.

In addition,  $\text{Ca}^{2+}$  is important for the effects of phospholipids and activity of relevant enzymes. The interaction of the PLD C2 domain with  $\text{Ca}^{2+}$  is stimulated by phospholipids [106], and  $\text{Ca}^{2+}$  sensitivities of different PLD isoforms vary from the micromolar to the millimolar range [10,107,108]. Increased  $[\text{Ca}^{2+}]_{\text{cyt}}$  evokes conformational changes in PLDs to assist with the recruitment of substrate  $\text{PtdIns}(4,5)\text{P}_2$  [109]. Additionally, enzymatic activity of PLC is regulated by  $\text{Ca}^{2+}$  through binding with the EF-hand motif of PLC [110], and PLA2s are classified as  $\text{Ca}^{2+}$ -dependent or -independent members [2].

### Cross-talk with hormone signalling

Auxin transporters is regulated by phosphorylation, and PINOID plays a fundamental role in the asymmetrical localization of membrane proteins during polar auxin transport by regulating the polarity of PIN's apical-basal localization [111]. Interestingly, PDK1 (phosphoinositide-dependent kinase 1) enhances PINOID activity and is involved in polar auxin transport regulation [111]. Both PDK1 and PINOID can bind to PA,  $\text{PtdIns}3\text{P}$ ,  $\text{PtdIns}(3,4)\text{P}_2$  and  $\text{PtdIns}(4,5)\text{P}_2$  through the PH (pleckstrin homology) domain [111,112]. PP2A (protein phosphatase 2A) and PINOID act

antagonistically on mediating PIN apical-basal polar localization [113] and PP2A can bind to PA [114]. In addition, PLD $\zeta$ 2 and product PA are required for the normal cycling of PIN2-containing vesicles to maintain auxin transport, and the deficiency of PLD $\zeta$ 2 results in suppressed sensitivity to auxin and reduced root gravitropism [20]. These findings suggest that membrane phospholipid turnover is important for regulating polar auxin transport (Figure 2).

Recently several studies revealed the crucial functions of phospholipids in vesicular trafficking. The trafficking of auxin efflux carrier PIN2 protein is dependent on SNX1 (sorting nexin 1)-containing endosomes which are sensitive to the PI3K inhibitor wortmannin [115]. SNX1 co-localizes with PtdIns3P-enriched membrane subdomains, and is a candidate downstream effector of PI3K because of the presence of phosphatidylinositol-binding PX (phox homology) domain of SNX1 [116]. During the regulation of turnover of auxin transporters, PIN2 is targeted to the vacuole for degradation, and the trafficking of PIN2-containing vacuolar depends on the phosphatidylinositol pathway [116]. Wortmannin affects the recycling of vacuolar sorting receptors between the PVC (prevacuolar compartment) and the TGN (*trans*-Golgi network) [116–118] and whose treatment inhibits the trafficking and degradation of PIN2, and increases the total PIN2 protein level in membrane [116]. Such cross-talk prompts us to ask questions. Do the membrane compositions affected by phospholipids influence the localization of membrane protein PINs? Do phospholipids serve as mediators to recruit the proteins to membrane sites and to promote signal transduction? Although the current studies suggest that the latter speculation may be the case, the relevant mechanisms are still unclear.

*At5PTase13* regulates the expression of *CYP83B1*, which is important for auxin homeostasis [40], and overexpression of *AtIPK2 $\beta$*  leads to more axillary shoot branching and an IAA-related phenotype by mediating the transcription of the IAA-related genes *CYP83B1*, *PIN4*, *MAX4* and *SPS* [95].

Only PA was reported to bind CTR1 (constitutive triple response 1), which negatively regulates ethylene responses in *Arabidopsis* [119] to disrupt the intramolecular interaction between the CTR1 kinase domain and the CTR1 N-terminal regulatory domain to inhibit its kinase activity [49] (Figure 2).

### Regulation through PA

PA, the simplest membrane phospholipid, has emerged as a new class of lipid mediator involved in various cellular processes, such as signal transduction, membrane trafficking, secretion and cytoskeletal rearrangement [10]. Cellular PA is generated mainly via two routes: DGK (DAG kinase) phosphorylating DAG or PLD hydrolysing structural phospholipids. In plants, PA serves as a second messenger and is triggered in response to various biotic and abiotic stresses, including pathogen infection [77,120], drought, salinity, wounding, cold, cell death [68,70,98] and oxylipin production [71–73]. The PA signal production is fast (minutes) and transient.

PA plays a role in the ABA response during stomatal movement and seed germination. PLD $\alpha$ 1-derived PA binds to ABI1, a negative regulator of the ABA response, and inhibits its activity to promote stomata closure [44]. T-DNA (transferred DNA) insertional mutants of PA catabolic enzyme LPP (lipid phosphate phosphatase), *lpp2-1* and *lpp2-2*, are hypersensitive to ABA during germination due to the increase in PA levels [121].

Regarding the roles of PA in the abiotic stress response, it is suggested that PA produced by high PLD $\alpha$ 1 activity destabilizes the membrane and increases membrane leakage, whereas a regulated increase of plasma membrane PA by PLD $\delta$  may produce

signalling PA species that mitigate stress damage [63,65]. The PA level increases during chilling and cold acclimatization, as well as in response to freezing [63,122], and the results of microarray analysis imply that PA produced from the PLD and PLC/DGK pathways are involved in two different pathways in cold responses [62]. Osmotic stress also induces the rapid accumulation of PA [60,123], which may be involved in regulation of proline biosynthesis [124]. Studies using soya bean cells show that PA can bind MAPK6-related protein, an important mediator in stress and ethylene signalling, suggesting a different mechanism of PA involvement in the stress response [125] (Figure 2). Furthermore, the PA-interacting/binding proteins should be studied in more detail to illustrate the downstream mechanisms.

### Regulation through cytoskeleton

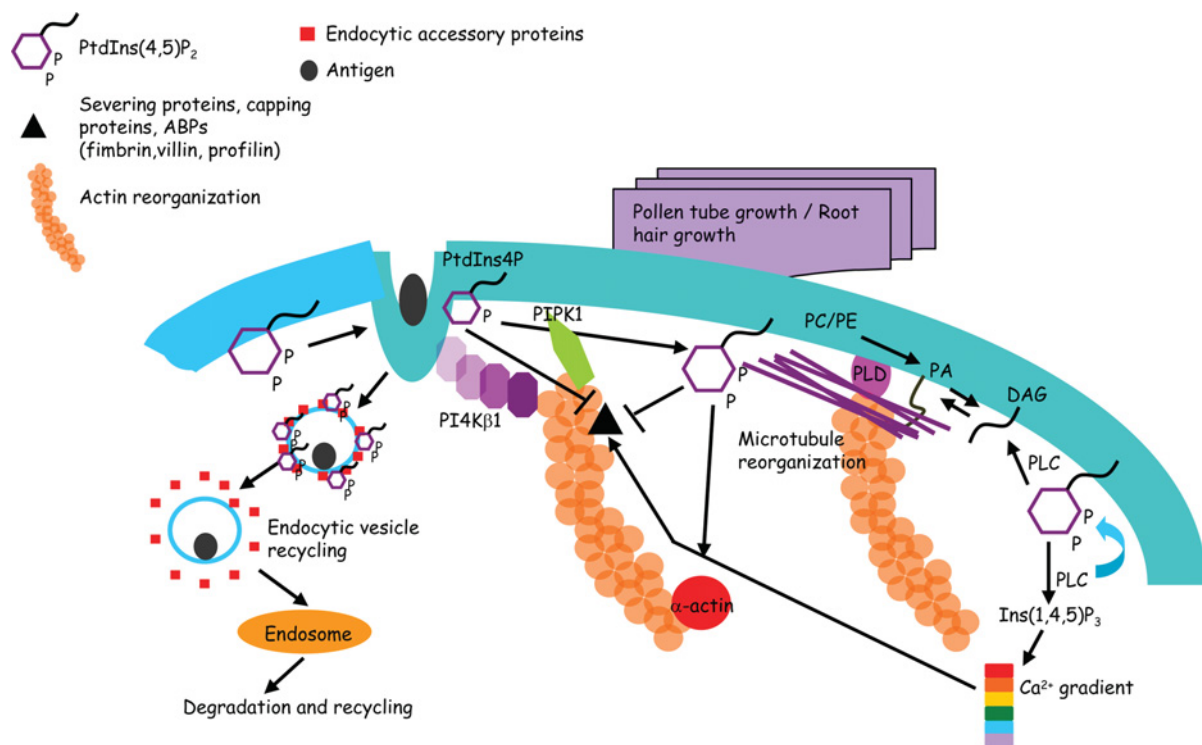
Both the actin and microtubular cytoskeletons can be regulated by phospholipids and participate in regulating the tip growth of the root hair and pollen tube. PtdIns(4,5) $P_2$ , Ins(1,4,5) $P_3$ , PA and the related PI3K, PI4K and PIP5K are involved in the organization of actin filaments; the microtubular cytoskeleton is mainly modulated by PLD.

Phospholipids play an integral role in regulating the structure and dynamics of the cytoskeleton through interaction between PtdIns(4,5) $P_2$  and many actin-binding proteins, including profilin, gelsolin,  $\alpha$ -actin, cofilin, filamin and vinculin [126–130]. It has been shown that PtdIns(4,5) $P_2$  controls the dynamic organization of F-actin (filamentous actin) through regulating profilin, a G-actin (globular actin)-binding protein [131], and actin remodelling in root hairs and pollen tubes, sites of rapid growth in plants, is sensitive to alterations of PtdIns(4,5) $P_2$  biosynthesis. Expression of mutant *Arabidopsis* Rac2 in tobacco pollen tubes decreases the plasma membrane PtdIns(4,5) $P_2$  and disrupts the normal actin filament orientation [132]. *Arabidopsis* PIP5K1, which synthesizes PtdIns(4,5) $P_2$  *in vivo*, interacts directly with actin and recruits PI4K $\beta$ 1 to the actin cytoskeleton [133]. Although it has not been confirmed, Stenzel et al. [30] speculated that the regulation of membrane-associated PIP5K3 in root hair elongation might relate to the control of F-actin (Figure 4).

The PLC-mediated PtdIns(4,5) $P_2$  turnover also affects the actin structure and normal pollen growth [38,134]. Elevation of intracellular PtdIns(4,5) $P_2$  [through photolysis of caged PtdIns(4,5) $P_2$ ] leads to the perturbation of apical morphology and the appearance in this region of a fine cortical mesh of microfilaments [23].

PA promotes actin polymerization by binding to and inhibiting the activity of AtCP, an actin-capping protein [135]. *Arabidopsis* PLD $\beta$ 1 can bind directly to actin, and the activity of PLD is, in turn, regulated by the actin cytoskeleton [136]. *AtSAC1* and *FRA3*, which encode PtdIns(3,5) $P_2$  phosphatase and 5PTase respectively, are both required for actin organization and cell wall synthesis [41,42]. Deficiency of *AtSfh1*, a P1TP-encoding gene, results in arrested root hair pattern formation by altering both actin and the microtubular cytoskeleton [24]. A recent study shows that both PtdIns3P and PtdIns4P modulate actin filament organization in guard cells of day flowers [137].

The involvement of the microtubular cytoskeleton in phospholipid signalling is mediated by PLD. Treatment with the PLD inhibitor butan-1-ol disrupts cortical microtubule organization in soya bean cells and fucoid alga [138,139] and promotes microtubule depolymerization and release from the plasma membrane in *Arabidopsis* [140]. Unlike in mammalian cells, cortical microtubules lie beneath the plasma membrane in plant cells, and PLD is postulated to be the linker [141].



**Figure 4** Phospholipids participate in the tip growth of the root hair and pollen tube through regulating actin and microtubule cytoskeletons

PtdIns(4,5) $P_2$  is synthesized on the plasma membrane and assembles a number of endocytic accessory proteins to induce the antigen-stimulated endocytosis. PIP5K1 and PtdIns(4,5) $P_2$  interact directly with actin and recruit PI4K $\beta$ 1 to actin. PtdIns(4,5) $P_2$  interacts and suppresses the activity of actin-binding proteins, including severing proteins, capping proteins, ABPs (actin-binding proteins: fimbrin, villin, profilin), and induces the activity of  $\alpha$ -actin to regulate actin remodelling. PA turnover from DAG is the substrate for PLD which is involved in microtubule reorganization. Ins(1,4,5) $P_3$ - $Ca^{2+}$  oscillation regulates the activity of actin-binding proteins to participate in actin remodelling. Abbreviations: PC, phosphatidylcholine; PE, phosphatidylethanolamine.

## Others

Other kinds of regulation of phospholipids and relevant key proteins also exist. PLD activity, with the exception of  $\zeta$ -class PLDs, is dependent on calcium, whereas  $\beta$ -,  $\gamma$ - and  $\zeta$ -class PLDs require PtdIns(4,5) $P_2$  for *in vitro* activity [142]. A recent study shows that aluminium ions inhibit PLD in a microtubule-dependent manner [143]. The activity of AtPIP5K1 is inhibited by phosphorylation through protein kinase A [144]. In addition, the MORN motifs in OsPIP5K1 (*Oryza sativa* PIP5K1) and the highly charged region of OsPI4K2 are involved in the regulation of subcellular localization and substrate binding [145,146]. The WD40 domain of 5PTase13 interacts with SnRK1 and acts as a positive regulator of SnRK1 activity to stabilize the protein and suppress its degradation [84].

## CONCLUSIONS AND PERSPECTIVES

Substantial evidence demonstrates that phospholipid signalling plays an important role in various aspects of plant growth and development, stress response and hormone signalling, and extensive progress has been made toward understanding the regulation and function of phospholipid molecules and related enzymes.

It has been noted that several gene families involved in phospholipid signalling contain more than ten isoforms in *Arabidopsis*, including PIP5K, PLD and 5PTase, and isoforms in the same family have distinct expression patterns and may have specific functions. The precise analysis of the expression patterns in specific tissues, at different developmental stages and under

different stimulations of individual isoforms will facilitate the dissection of their specific functions.

Although post-transcriptional regulation is an important process in functional modulation, studies of the regulation of phospholipid-related enzymes and protein level are limited. PIP5K activity can be inhibited by phosphorylation and modulated by changes in subcellular localization and substrate binding [145,146], shedding light on the cellular regulation of phospholipid signalling.

The IP $_3$ R [Ins(1,4,5) $P_3$  receptors] have been identified from animal and yeast cells; however, none are known in plants. Ins(1,4,5) $P_3$  activates IP $_3$ R in the endoplasmic reticulum, resulting in the  $Ca^{2+}$  release and leading to the  $[Ca^{2+}]_i$  oscillations. In plants, an external  $Ca^{2+}$ -sensing receptor (CAS) serves as a receptor to regulate Ins(1,4,5) $P_3$  levels and then in turn trigger the  $Ca^{2+}$  release. Although cell-surface receptors and IP $_3$ R are unknown [2,83], it is reasonable to speculate that CAS may also act as a candidate receptor for Ins(1,4,5) $P_3$  after studies using biochemical and single-cell imaging analyses [83]. RACK1 (receptor for activated C-kinase 1), which has highly conserved WD40 repeats, acts as a scaffold protein through interacting with IP $_3$ R and other proteins in animal nervous systems [147]. In *Arabidopsis*, the first crystal structure of RACK1 isoform A has been characterized [148]. Does the WD40 domain of 5PTase family also play a role as a scaffold protein for IP $_3$ R in plants? In addition, the functions of Ins(1,4,5) $P_3$ -mediated  $Ca^{2+}$  oscillation seems to be more important than Ins(1,4,5) $P_3$  itself in *Arabidopsis*; do Ins(1,4,5) $P_3$  and  $Ca^{2+}$  share the same receptor?

Until now, most studies have been performed using the dicotyledonous model plant *Arabidopsis*; few studies have been carried



out in rice [15,149]. Whether the function is conserved in monocotyledons (such as rice) is not well understood. Preliminary analysis reveals a difference in the domain structure and isoform numbers of key enzymes in the PLD family [15], suggesting a functional divergence between dicotyledons and monocotyledons. Understanding these differences may provide further information for agricultural improvement.

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