The Arabidopsis thaliana Immunophilin ROF1 Directly Interacts with PI(3)P and PI(3,5)P and Affects Germination under Osmotic Stress

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Abstract

A direct interaction of the Arabidopsis thalianaimmunophilin ROF1 with phosphatidylinositol-3-phosphate and phosphatidylinositol-3,5-bisphosphate was identified using a phosphatidylinositol-phosphate affinity chromatography of cell suspension extracts, combined with a mass spectrometry (nano LC ESI-MS/MS) analysis. The first FK506 binding domain was shown sufficient to bind to both phosphatidylinositol-phosphate stereoisomers. GFP-tagged ROF1 under the control of a 35S promoter was localised in the cytoplasm and the cell periphery **M**icotiana tabacum leaf explants. Immunofluorescence microscopy of Arabidopsis thalianaroot tips verified its cytoplasmic localization and membrane association and showed ROF1 localization in the elongation zone which was expanded to the meristematic zone in plants grown on high salt media. Endogenous ROF1 was shown to accumulate in response to high salt treatmerAntabidopsis thaliana young leaves as well as in seedlings germinated on high salt media (0.15 and 0.2 M NaCl) at both an mRNA and protein level. Plants over-expressing ROF1, (WSROF10E), exhibited enhanced germination under salinity stress which was significantly reduced in therof1² knock out mutants and abolished in the double mutants of ROF1 and of its interacting homologue ROF2 (WSrof12²). Our results show that ROF1 plays an important role in the osmotic/salt stress responses of germinating Arabidopsis thaliana seedlings and suggest its involvementin salinity stress responses through a phosphatidylinositol-phosphate related protein quality control pathway.

Citation: Karali D, Oxley D, Runions J, Ktistakis N, Farmaki T (2012) ATabe dopsis thaliandmmunophilin ROF1 Directly Interacts with PI(3)P and PI(325) Rd Affects Germination under Osmotic Stress. PLoS ONE 7(11): e48241. doi:10.1371/journal.pone.0048241

Editor: Haibing Yang, Purdue University, United States of America

Received February 17, 2012Accepted September 21, 2012Published November 2, 2012

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Funding: This research project is co-financed by the E.U.-European Social Fund (75%) and the Greek Ministry of Development-GSRT (25%) (PENED) as well as a European Molecular Biology Organization (EMBO) short term fellowship ASTF 395.00-2007 and the Biotechnology and Biological Sciences Resemble C (BBSRC). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing Interests: The authors have declared that no competing interests exist.

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Introduction

Phosphatidylinositol phosphates (PIPs) are phospholipids with important regulatory properties in signaling and trafficking processes [1]. PIPs possess a characteristic subcellular localizatimses ucselor52inllular thelonth7egucognion propeler structure may be utilized for binding [20]. Because of

Bacterially Expressed ROF1 Interacts with PI(3)P and PI(3,5)P through its FKBD Domains

amount of protein remained inside the inclusion bodies. The results suggest a differential capacity of the FKBD domains for

ROF1 and its truncated mutants (Figure 2B; Table S1; Table lipid binding as well as the ability of ROF1 to bind to PI(3,5))P S2) were over-expressed with an N-terminal GST tag and Cthe absence of the FKBD1 inspite the fact that FKBD2 and terminal-His tag following transformation of a BL21 bacterial cell FKBD3 do not possess a structural identity (compared to the same extent as the line with the pALEX vector used for their cloning. Following predictable PIP binding sites, at least not to the same extent as the double purification using Glutathione and Nickel columns ROF1 FKBD1 does (Figure S2).

and its truncated mutants were eluted and used in a lipid overlay

assay (Figure 3). ROF1 specifically associated with both the PI(3) be of the $Av\tau\iota$ -ROF1 for the Characterization of the as well as with the PI(3,5) (Figure 3A). The FKBD1 domain was PI(3,5) P Binding Site

shown sufficient for PIP binding and appeared to possess a higher A polyclonal antibody was raised against the bacterially affinity for the PI(3,5)Pstereoisomer and reduced affinity for the expressed ROF1 as described in the Materials and Methods. PI(3)P (Figure 3A, B). A competition assay, using the two Taking into account that αντι-ROF1 was raised against the stereoisomers as competing lipids, produced a specific reduction the account that αντι-ROF1 was raised against the stereoisomers as competing lipids, produced a specific reduction the BL21 cells were purified and run on a polyacrylamide gel. a competing lipid reduced signal for both the PI(3)P and the Western blotting using

PI(3,5)^B in the ROF1 and the FKBD1 assay (Figure 3A, B). Although the FKBD1 was shown to possess a high affinity for the

PI(3,5)B its absence did not abolish binding to the $PI(3_25)P$

stereoisomer as demonstrated by the ability of the 2FKTPR

construct to interact with both lipids (Figure 3C). On the other

hand, FK3TPR, only interacted with the PI(3)P (Figure 3C). Although the TPR-calmodulin binding domain alone was also

over-expressed, we were unable to solubilise it and the entire

Figure 3. Lipid overlay assays in order to characterize ROF1 binding to different PIP stereoisomers. Dot blotting of different PIP stereoisomers probed with 1µg/ml of the ROF1 protein (A) and its truncated mutant 1FK (FKBD1) (B) (Figure 2B; Figure S1) and detected with GST. 2): No competing lipid has been added in the incubation, [PI(3,5]P50µM of the lipid has been added during the incubation; [PI(3)P]: 50M of the lipid has been added during the incubation. (C) Dot blotting of different PIP stereoisomers probed withµg/ml of the ROF1 truncated mutants 2FKTPR and FK3TPR (Figure 2B; Figure S1) and detected withGST. (D) Characterization of the binding site of ROF1 to PI(3)P and PI(3,5)P using the N-terminus specific antibody anti-ROF1. Overexpressed ROF1 was pre-incubated with different concentrations of the antibody, as indicated, prior to its use in a lipid overlay assay and detected with anti-GST. doi:10.1371/journal.pone.0048241.g003

showed a complete recognizing ability for the ROF1 providing differences were, however, observed in ROF1 protein accumulathat the polylysine rich patch (KKKLLK) was present (NROF1 tion at high salt concentrations (1 M) compared to lower salt construct) (Figure 2B; Figure S1; Figure S3C). Taking into account oncentrations (0.2 M) but an earlier reduction was observed for the specific site of antibody recognition, we used the antibody inthe 1 hr 1 M NaCl treatment compared to the 0.2 M incubation. order to block ROF1 prior to the incubation in the lipid overlay Protein accumulation was also observed in the case of whole assay. Different concentrations of the ROF1 antibody were used iseedlings, germinated in high salt MS media (0.15 and 0.2 M a mixture with the protein (0 to µl/ml) and following incubation NaCl) compared to the control seedlings (Figure 5D). The in a lipid overlay assay as previously described 4-GST was used for the detection of protein-lipid interaction (Figure 3D). Theonly 3 hrs after water imbibition (Figure 5E). antibody pre-incubation reduced binding affinity for the 3,5

phosphorylated lipid but not for the lipid phosphorylated at the Over-expressed and Endogenous ROF1 Localization third position of the inositol ring (Figure 3D). Under Control and Osmotic/salinity Stress Conditions

Antibody specificity was further verified using both thaliana wild type and mutant lines (WDsLOX, W6ff, WSoff/2²) stressed at 3°C for up to 4 hrs. Results showed (Figure 4; Figure ROF1. A cytoplasmic localization of the protein was observed S4) that the xvti-ROF1 specifically recognizes endogenous ROF1 using a 3D reconstruction (Figure 6A) as demonstrated by the and to a very low extent its homologous partner ROF2 providing traces (empty spaces) that the embedded organelles mark in the the latter has been accumulated following heat stress [24] tytoplasm and the cytoplasmic strands formed. A peripheral localization may also be observed, more clearly seen in a single

localization may also be observed, more clearly seen in a single confocal section (Figure 6B). ROF1 localization was distinct to GFP localization since no nuclear localization of the GFP tagged-

ROF1 is Induced Under Salinity Stress

3 week oldA. thalianplants were incubated with 0, 0.2 or 1 M ROF1 was observed (Figure 6C). In order to confirm the nature of NaCl. In young leaves collected at different time points, ROF1 the peripheral localization of ROF1, a plasmolysis effect caused by and ROF2 accumulation was observed at an mRNA level 0.25 M NaCl treatment was used. As it is shown (movie S1), the following salt treatment (Figure 5A). A fast induction within peripheral staining of ROF1 is not due to a protein association 3 mins appeared following incubation with 1 M NaCl for both with the cell wall.

ROF homologues. mRNA accumulation was also observed in the In order to further verify ROF1 subcellular localization case of ROF1 in seedlings germinated on high salt/osmotic mediammunofluorescence microscopy with thalianaroot tips was compared to the control (MS media). However, this accumulationused.A. thalianareedlings were germinated on 0.8% agar media was not observed for ROF2 (Figure 5B). ROF1 accumulation wator wator wator water a protein level (Figure 5C). No apparent performed as described in the Materials and Methods. Incubation

Figure 4. Recognition of ROF1 and ROF2 using anti-ROF1. 5-day old A. thalianawild type, ROF1 knock-outr ϕ f1²) of both Columbia and Wassilevskija background and ROF1/2 double mutants (M/f5²/2²) were incubated at 37C for up to 4 hrs. anti-ROF1 specifically detects the ROF1 protein in plant extracts and to a less extent a second upper band (ROF2) following heat treatment only. Columbia ROF1 knock-out belongs to the WDsLox line. doi:10.1371/journal.pone.0048241.g004

with pre-immune serum oxvtt-ROF1 staining of of f plants did changed, expanding throughout the meristematic zone (Figure 7C; not produce any fluorescence (results not shown) confirming lovie S3). The cytoplasmic localization of ROF1 was confirmed antibody specificity as described above. In untreated plants and/ith immunofluorescence microscopy (Figure 7A; Movie S2) plants treated for a short period of time with high NaCl (0.4 M for whereas a stronger peripheral staining was observed in plants 2 mins) ROF1 was mainly localized in the elongation zone of thegrown on high salt media (Figure 7C; Movie S3) probably root tip (seedlings were immersed in the solution washed brieflyndicating membrane association. In any case, no nuclear and then fixed) (Figure 7A, B; Movie S2). However, in plantslocalization was observed.

grown on high salt media (0.2 M NaCl), the localization pattern

Figure 5. Gene expression and protein accumulation study for ROF1. (A) Real time PCR of wild typA. thalianaplants treated for different time points and under different salt concentrations (0 M, 0.2M and 1M NaCl). Results are expressed as fold increase relative to the control set at 1. Control represents mock treatment with ^I₂O for the respective time points to the NaCl treatment. (B) Real time PCR of control (C) wild type 4-day old A. thalianaseedlings germinated on MS media and of seedlings germinated and grown on media containing NaCl or mannitol. Results are expressed as fold increase relative to the control set at 1. Control represents plants grown on MS media. Results are average of three independent experiments performed in triplicates. (C) Western blotting of wild typeA. thalianayoung leaves obtained from plants which have been treated for different time points and at different salt concentrations (0 M, 0.2M and 1M NaCl). Control represents mock treatment with ^I₂O for the respective time points to the NaCl treatment. (B) Real time PCR of seeds 3 hrs after water imbibition. doi:10.1371/journal.pone.0048241.g005

Figure 6. ROF1 localisation using tobacco plants transiently transformed and expressing GFP tagged ROF1 and GFP only. (A) Control

Effect of Osmotic/ionic Stress on the Germination of Wild Type and ROF MutantA. thalianaSeeds Several lines of Columbia wild type (Col-WT), Columbia

Several lines of Columbia wild type (Col-WT), Columbia f (Col-rof²), Wassilevskija wild type (WS), WG² and WSof²/ concentrations. Although no difference was recorded on the rof², WS complemented mutants (WSROF1CM) and WS ROF1 germination rate of plants grown on media without NaCl in over-expressions (WSROF10E) were confirmed by western⁶elation to the WT (Figure S5A, D) a significant delay was

recorded on the germination ouff plants germinated on media ROF1 AffectsA. thaliana Seed Germination through containing salt concentrations of 0.2 M NaCl (Figure S5B). Thea Phosphatidyl-Inositol-3 Kinase (PI3K) Related Pathway NaCl effect on plant germination was also observed with the A. thalianaWSROF10E and WSoff /2² seeds were germi-Wassilevskija wild type another (Figure 8A). Germination was nated in the presence or absence of wortmannin or LY294002 nearly eliminated at 0.2 M NaCl when the $VMSF^2/2^2$ double under both control and salinity stress conditions. Germination mutants were used (Figure 8A, C). In order to show that the osmotic effect of ROF1 is exclusively due to the loss of the ROF1 gene complemented Worf² as well as WSROF10E lines were used. It was shown that gene complementation completely recovered the wild type phenotype whereas gene over-expression resulted in a highly enhanced germination under salinity stress (Figure 8A). In order to differentiate between the ionic and the osmotic effect, WS, WSROF1OE, WtSft², WSROF1CM, WSrof² and WSroft²/ 2^2 were geminated on media containing 0.4 M mannitol and the osmotic effect on plant germination was confirmed (Figure 8B, C; Figure S5C).

Figure 8. Germination efficiency of ROF1 and ROF2 mutants under osmotic/salinity stress. (A) Germination rate of the Wassilevskija background plants WS, $W\mathfrak{B}1^2$, $W\mathfrak{S}of2^2$, $W\mathfrak{S}of1^2/2^2$, $W\mathfrak{S}OF1CM$ and $W\mathfrak{S}ROF1OE$ on MS media containing 0.2 M NaCl. (B) Germination rate of the wild type Wassilevskija background plants WS, $W\mathfrak{B}1^2$, $W\mathfrak{S}of2^2$, $W\mathfrak{S}of1^2/2^2$, and $W\mathfrak{S}ROF1OE$ on MS media containing 0.4 M mannitol. Three independent characterized lines (in each experiment) were used for each genotype. The results show the average of three independent experiments

Figure 9. Effect of PI3K inhibitors on seed germination. Seed germination efficiency of WSROF1OE and $MS^2/2^2$ at different time points under control (A) or salinity stress (0.1 M NaCl) (B) in the presence or absence qIM3 wortmannin or 60 μ M LY294002. The results show the average of four independent experiments performed for each treatment. Values are means and bars are SDs. doi:10.1371/journal.pone.0048241.g009

our work suggests its involvement in salt stress tolerance since its absence significantly reduces and further abrogates germination efficiency under salinity stress in the $f^2/2^2$ double mutant (Figure 8). Unlike ROF1, ROF2 mRNA and protein do not accumulate-at least not to a detectable by our antibody levelneither under control (also shown by previous studies) nor under salinity stress (as it occurs during heat stress) (Fig 5). ROF2 became detectable only following its enrichment specifically on the PI(3,5)B column as demonstrated by our mass spectrometry data, pointing to its specific involvement in a PIP associated osmotic stress related mechanism. The involvement of such a mechanism in osmotic/salinity stress responses was supported by experiments with the PI3K inhibitors wortmannin and LY294002 (Figure 9). Our results point to a PI3K related mechanism involved in seed germination under salinity stress through a ROF1 function. Seeds over-expressing ROF1 showed better germination efficiency in the presence of the inhibitors under normal conditions compared to the knock out plants. In contrast and under salt stress conditions and inhibitor addition, ROF1 overexpression did not enhance seed germination (as it happens in the absence of the inhibitor) and

Fluorescence and Confocal Microscopy

Localization studies of GFP-tagged ROF1 as well as immunofluorescence microscopy were performed using fluorescence and Confocal microscopy. Fluorescence microscopy was performed using an Axioskop 40 fluorescence microscope with a A-Plan Varel contrast 40x/0.65 (Zeiss, Germany). Images were captured using a ProgRes digital camera system from Jenoptic (Germany). For confocal microscopy two confocal microscopes were used; Live cell imaging was performed using a laser Scanning confocal Microscope TCS SP5 SN: 5100000134 with an HCX PL APO CS 63.0×1.20 water UV lens. A pinhole 1 and 1.6 zoom were used. Scan speed was 400 Hz. Resolution was 102024 with 8 bits/ pixel. Excitation was achieved with a 488 Argon laser for the GFP and a DPSS 561 for the RFP. Emission bandwidth for GFP was 504.0 nm-521.2 nm and the emission bandwidth for RFP 574.2 nm - 620.1 nm. Fixed tissue imaging was performed using an HC X PLAPO CS 40.0×0.85 dry lens. Pinhole was 2, zoom 1.7 and resolution 10241024. Excitation was achieved using a 405 nm Diode for DAPI and a 561 DPSS for Cv3 red fluorescence. Emission bandwidth for DAPI was 423 nm -475.6 nm and the emission bandwidth for Cv3 567 nm -619.4 nm. Image analysis was performed using the LAS AF software version 2.1.0 as well as the ImageJ (Java) software. Also, a Zeiss LSM 510 confocal microscope with a EC Plan-Neofluar 40x/1.30 oil DIC M27 objective was used. was 26 fluorescence4

 $\frac{1}{(AVI)} \xrightarrow{f_1} \frac{m_1}{f_1} \xrightarrow{f_1} \frac{m_2}{f_1} \xrightarrow{f_1} \frac{1-C}{f_1} \xrightarrow{f_1} \frac{1-C}{f_1} \xrightarrow{f_1} \frac{1}{f_1} \xrightarrow$

Acknowledgments

We would like to thank Prof. Adina Breiman for the Wassilevskija seeds, WS, WSroff , WSroff , WS ,W28

- Farmaki T, Sanmartin M, Jimenez P, Paneque M, Sanz C, et al. (2007) Differential distribution of the lipoxygenase pathway enzymes within potato chloroplasts. J Exp Bot 58: 555–568.
 Zou X, Osborn MJ, Bolland DJ, Smith JA, Corcos D, et al. (2007) Heavy chain
- Zou X, Osborn MJ, Bolland DJ, Smith JA, Corcos D, et al. (2007) Heavy chain only antibodies are spontaneously produced in light chain-deficient mice. J Exp Med 204: 3271–3283.
- Panagiotidis CA, Silverstein SJ (1995) pALEX, a dual-tag prokaryotic expression vector for the purification of full-length proteins. Gene 164: 45–47.
- 49. Kargiotidou A, Deli D, Galanopoulou D, Tsaftaris A, Farmaki T (2008) Low temperature and light regulate delta 12 fatty acid desaturases (FAD2) at a(ts)af\$sciptional level in cotton (Gossypium hirsutum Td [((potat2i1287(7Pal.(13rTJ -10(Me9:t7io09(in5(5)406cFAD294.892 11.3976 Td [509.)-717(Kargiotido)-14(u)2692(A,)276

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