

1 PLASMA-MEMBRANE H<sup>+</sup>-ATPase-DEPENDENT CITRATE EXUDATION FROM  
2 CLUSTER ROOTS OF PHOSPHATE-DEFICIENT WHITE LUPIN

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32 RUNNING TITLE: PM H<sup>+</sup>-ATPase-dependency of citrate exudation

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34

35 ABSTRACT

36 White lupin (*Lupinus albus* L.) is able to grow on soils with sparingly available phosphate (P) by  
37 producing specialized structures called cluster roots. To mobilize sparingly soluble P forms in  
38 soils, cluster roots release substantial amounts of carboxylates and concomitantly acidify the  
39 rhizosphere. The relationship between acidification and carboxylate exudation is still largely  
40 unknown. In the present work, we studied the linkage between organic acids (malate and citrate)  
41 and proton exudation in cluster roots of P-deficient white lupin. After the illumination started,  
42 citrate exudation increased transiently and reached a maximum after 5 hours. This effect was  
43 accompanied by a strong acidification of the external medium and alkalization of the cytosol,  
44 as evidenced by *in vivo* NMR analysis. Fusicoccin, an activator of the plasma-membrane H<sup>+</sup>-  
45 ATPase, stimulated citrate exudation, whereas vanadate, an inhibitor of the H<sup>+</sup>-ATPase, reduced  
46 citrate exudation. The burst of citrate exudation was associated with an increase in expression of

47 the *LHA1* plasma-membrane H<sup>+</sup>-ATPase gene, an increased amount of H<sup>+</sup>-ATPase protein, a  
48 shift in pH optimum of the enzyme and post-translational modification of an H<sup>+</sup>-ATPase protein  
49 involving binding of activating 14-3-3 protein. Taken together, our results indicate a close link in  
50 cluster roots of P-deficient white lupin between the burst of citrate exudation and plasma-  
51 membrane H<sup>+</sup>-ATPase catalyzed proton efflux.

52

53 KEYWORDS:

54 *Lupinus albus*, organic acids, root exudates, malate, rhizosphere, proton pump, phosphorus,  
55 carboxylate release, pH

56

57 INTRODUCTION

58

59 The availability of phosphate (P) is a wide-spread limiting factor for plant growth (Handreck  
60 1997; Marschner 1995). To deal with this situation, plants have developed several strategies to  
61 improve P acquisition. The most common one is the association with mycorrhizal fungi. If plants  
62 do not engage in symbiosis, they adapt to phosphate-poor soils by increasing the quantity and  
63 length of roots and root hairs, and in many cases by increasing the synthesis and the release of P-  
64 mobilizing root exudates, such as organic acids, phenolics, and acid phosphatases (Dinkelaker,  
65 Römheld & Marschner 1989; Neumann et al. 2000; Tomasi et al. 2008).

66 Some plants have developed special root structures in response to phosphate starvation, such as  
67 the dauciform roots found in the Cyperaceae family (Davies, Briarty & Rieley 1973; Lamont  
68 1974; Shane, Dixon & Lambers 2005) and the cluster or proteoid roots of many Proteaceae  
69 (Purnell 1960). Cluster roots or proteoid roots are generally bottlebrush-like root structures.  
70 Many studies have demonstrated the capacity of these roots to increase the availability of soil  
71 phosphate (Dinkelaker et al. 1989; Neumann et al. 1999; Shane et al. 2003). Phosphate  
72 mobilization is suggested to occur predominantly via the release of large amounts of organic acid  
73 anions by the cluster roots (Dinkelaker et al. 1989; Gardner, Parbery & Barber 1982; Jones &  
74 Darrah 1994; Lambers et al. 2002).

75 White lupin (*Lupinus albus* L.) has often been used as a model plant for studying P-deficiency  
76 response and cluster-root formation (Neumann & Martinoia 2002). In cluster roots of white  
77 lupin, carboxylate release follows a spatial and temporal pattern. In the first growing phase,  
78 cluster roots release low amounts of organic acids, mainly malate. In mature proteoid roots, a  
79 burst of citrate exudation can be observed which is accompanied by acidification of the

80 rhizosphere (Neumann et al., 2000). Citrate is likely to be exuded via an anion channel, since  
81 exudation can be inhibited by anthracene-9-carboxylic acid (Neumann et al. 1999). In a more  
82 recent study, a citrate-specific channel from lupin root protoplasts was described (Zhang, Ryan  
83 & Tyerman 2004) which could be responsible for massive citrate exudation; however, regulatory  
84 mechanisms of carboxylate exudation still need to be elucidated. A correlation between citrate  
85 and proton extrusion has been suggested by several authors (Ligaba et al. 2004; Ohno, Koyama  
86 & Hara 2003; Sas, Rengel & Tang 2001; Shen et al. 2005; Zhu et al. 2005); however, in these  
87 studies different or even contrasting results were obtained, possibly due to the use of different  
88 experimental approaches and culture conditions.

89 The involvement of the plasma-membrane (PM) H<sup>+</sup>-ATPase in organic anion release via proton  
90 extrusion in white lupin was suggested by Kania et al. (2001) and Yan et al. (2002). Using  
91 isolated plasma-membrane vesicles from whole proteoid roots and lateral roots grown in the  
92 presence or absence of phosphate, these authors showed that proteoid roots exhibited an  
93 increased PM H<sup>+</sup>-ATPase activity that was correlated to changes in the amount of the  
94 corresponding protein. Ligaba et al. (2004) also observed an enhanced release of citrate, but not  
95 of malate, associated with the increased activity of the plasma-membrane proton pump in cluster  
96 roots of P-deficient *Lupinus pilosus*. Recently Shen et al. (2005), who investigated citrate  
97 exudation in aluminum-stressed soybean, demonstrated that citrate exudation was coupled to  
98 changes in the activity of the PM H<sup>+</sup>-ATPase, involving transcriptional and post-translational  
99 modification of the enzyme. Furthermore it has been reported that release of carboxylates from  
100 cluster roots of P-deficient white lupin may show diurnal variations (Watt & Evans 1999).

101 The goal of this work was to study the linkage between organic acids (malate and citrate) and  
102 proton exudation in fully grown cluster roots of P-deficient white lupin, highlighting the  
103 involvement of PM H<sup>+</sup>-ATPase activity and the regulatory aspects.

104

105

## 106 MATERIAL AND METHODS

107

### 108 *Plant material and growth conditions*

109 White lupin plants (*Lupinus albus* L. cv. Amiga, Südwestdeutsche Saatzucht, Rastatt, Germany)  
110 were grown in hydroponic conditions as described by Massonneau et al. (2001), with the  
111 exception that 48 plants were grown in 50-liter containers. Seeds were soaked overnight in  
112 aerated water, and then kept for 3 days in the dark followed by one day in the light, on filter  
113 paper soaked in 0.2 mM CaCl<sub>2</sub> to allow them to germinate. Seedlings were transferred to a  
114 hydroponic culture medium (0.05 mM Fe(III)-EDTA, 2.5 mM Ca(NO<sub>3</sub>)<sub>2</sub>, 0.9 mM K<sub>2</sub>SO<sub>4</sub>, 0.8  
115 mM MgSO<sub>4</sub>, 38 μM H<sub>3</sub>BO<sub>3</sub>, 12.5 μM MnSO<sub>4</sub>, 1.25 μM CuSO<sub>4</sub>, 1.25 μM ZnSO<sub>4</sub>, 0.33 μM  
116 (NH<sub>4</sub>)<sub>6</sub>Mo<sub>7</sub>O<sub>24</sub>, 62.5 μM KCl and with 0.25 mM KH<sub>2</sub>PO<sub>4</sub> in case of P-sufficient condition).  
117 Plants were grown at 22°C and 65% relative humidity with a light period of 16 h at 200 μmol m<sup>-2</sup>  
118 s<sup>-1</sup> for 5 weeks. For most of the experiments, the root samples were collected from P-deficient  
119 white lupins at different times after the start of the light (HALS). Only the fully grown parts of  
120 cluster root were collected (see also Fig. 1), corresponding to the immature and mature parts  
121 described by Massonneau et al. (2001). For the measurement of the cytosolic and vacuolar pH  
122 under -P and +P condition, we harvested the apical part (the last centimeter) of lateral roots.

123

124 *Quantification of citrate and malate released by cluster roots*

125 Collection of root exudates was performed according to Massonneau et al. (2001). Fully grown  
126 cluster roots were collected, pre-incubated in 10 mM CaSO<sub>4</sub> and incubated in 10 mM CaSO<sub>4</sub> in  
127 2-(N-morpholino) ethanesulfonic acid and 1,3-bis[tris(hydroxymethyl)-methylamino] (MES-  
128 BTP) 15 mM pH 6 in the absence or presence of 10 μM fusicoccin or 500 μM vanadate for one  
129 hour at room temperature with agitation. Root weight was determined and roots were used for  
130 the detection of root-induced change of pH (see below). Sample solutions were frozen at -80°C  
131 until processing. Citrate or malate content in the exudates was determined using the kit for citric  
132 acid or malic acid test (Boehringer, Mannheim, Germany) according to the manufacturer's  
133 instructions. Each dataset represents the mean of 3 independent experiments run in triplicates,  
134 processed and analyzed statistically (t-test, N=3, P<0.05) using Sigma Plot 8 (Systat software,  
135 Point Richmond, USA).

136

137 *Detection of root-induced change of rhizospheric pH*

138 The acidification capability of cluster roots was determined as described by Massonneau et al.  
139 (2001). After collection of the exudates, roots were placed on a glass plate covered by a 2-mm  
140 layer of 1% agar containing 0.04% (w/v) bromocresol purple at pH 6.0. Roots, which strongly  
141 acidify the medium, induce a color shift of the pH indicator from purple to yellow,  
142 corresponding to a pH below 5.5. Images were captured after 30 minutes of incubation.

143

144 *NMR measurements and P concentration*

145 Roots were collected and equilibrated for 10 min in the perfusion medium (0.5 mM CaSO<sub>4</sub>, 1  
146 mM MES-BTP, pH 6.1). <sup>31</sup>P-NMR spectra were recorded on a standard broad-band 10-mm

147 probe on a Bruker AMX 600 spectrometer (Bruker Analytische Messtechnik GmbH,  
148 Rheinstetten-Forchheim, Germany) equipped with Indy computer running XWIN-NMR version  
149 2.6. *In vivo* experiments were carried out by packing excised root segments in a 10 mm diameter  
150 NMR tube equipped with a perfusion system connected to a peristaltic pump in which the  
151 aerated, thermo-regulated (26°C) medium [0.5 mM CaSO<sub>4</sub>, 1 mM MES-BTP (pH 6.1)] flowed at  
152 10 ml min<sup>-1</sup>.

153 <sup>31</sup>P-NMR spectra were recorded at 242.9 MHz without lock, with a waltz-based broad-band  
154 proton decoupling and a spectral window of 16 kHz. Chemical shifts were measured relative to  
155 the signal from a glass capillary containing 33 mM MDP, which is at 18.5 ppm relative to the  
156 signal from 85% H<sub>3</sub>PO<sub>4</sub>. The spectra were determined using a 90° pulse angle and fast  
157 acquisition conditions (1<sup>s</sup> recycle time). Resonance assignments were performed according to  
158 Roberts et al. (1980) and Kime et al. (1982). Intracellular pH values were calculated from the  
159 chemical shift of the cytosolic and vacuolar Pi resonance after construction of a standard titration  
160 curve (Roberts et al. 1981).

161

#### 162 *Isolation of plasma membranes*

163 Plasma-membrane vesicles were isolated from fully grown cluster roots as following Santi et al.  
164 (1995) and Fisher-Schliebs et al. (1994). Briefly, 5 g of roots were homogenized with a mortar  
165 and pestle in a freshly prepared ice-cold extraction medium (250 mM sucrose, 2 mM MgSO<sub>4</sub>, 2  
166 mM ATP, 10% (v/v) glycerol, 10 mM glycerol-1-phosphate, 0.16% (w/v) BSA, 2 mM EGTA, 2  
167 mM DTT, 5.7% (w/v) choline-iodide, 1 mM PMSF, 20 µg ml<sup>-1</sup> chymostatin, 10 nM okadaic acid  
168 and 25 mM MES-BTP pH 7.6). Four milliliters of medium per g fresh weight of root tissue were  
169 used.

170 The homogenates were filtered through four layers of cheesecloth and the suspensions were  
171 subjected to differential centrifugation steps at 2°C: 1,500 g for 5 min (pellets discarded); 9,800  
172 g for 20 min (pellets discarded); 83,400 g for 30 min (pellets recovered); and 83,400 g for a  
173 further 30 min. Microsomes, gently resuspended in 1.2 mL of homogenization medium, were  
174 loaded onto a discontinuous sucrose gradient made by layering 2 mL of sucrose solution (1.13 g  
175 cm<sup>-3</sup>) onto a 3 mL sucrose (1.17 g cm<sup>-3</sup>) cushion, and centrifuged at 107600g for 2 h. The sucrose  
176 solutions were prepared in 5 mM MES-BTP, pH 7.4, and contained all of the protectants present  
177 in the homogenization medium. Vesicles migrating to the 1.13/1.17 g cm<sup>-3</sup> interface were  
178 collected, diluted with homogenization medium, and centrifuged at 122,400 g for 30 min. The  
179 pellets were resuspended in a medium containing 250 mM sucrose, 10 % (v/v) glycerol, 1 mM  
180 DTT, 50 µg mL<sup>-1</sup> chymostatin, 10 nM okadaic acid and 2 mM MES-BTP pH 7.0, were  
181 immediately frozen in liquid nitrogen, and stored at -80°C until use.

182

### 183 *Measurement of PM H<sup>+</sup>-ATPase activity and membrane protein content*

184 Plasma-membrane H<sup>+</sup>-ATPase activity was measured at 38°C in a 0.6 mL reaction (50 mM  
185 MES-BTP pH 6.5 or pH 6.2 to 8.0 for the pH dependency assay, 5 mM MgSO<sub>4</sub>, 100 mM KNO<sub>3</sub>,  
186 600 µM Na<sub>2</sub>MoO<sub>4</sub>, 1.5 mM NaN<sub>3</sub>, 5 mM ATP-BTP pH 6.5, 0.01 % (w/v) Brij 58  
187 (polyoxyethylene 20 cetyl ether), plus or minus 100 µM V<sub>2</sub>O<sub>5</sub>; the vanadate-dependent activity  
188 was 85±3%). The reaction was started by addition of membrane vesicles containing 0.5 µg of  
189 total protein; after 30 min, the reaction was stopped and color developed as previously described  
190 by Santi et al. (1995). Inorganic phosphate was quantified spectrophotometrically at 705 nm as  
191 described by Forbush (1983).

192 Protein content was determined as described by Bradford (1976), using BSA as standard, after  
193 solubilizing membrane vesicles with 0.5 M NaOH (Gogstad & Krutnes 1982).

194

#### 195 *Western blots*

196 Equal amounts of protein isolated at the different time points were loaded, electrophoresed in an  
197 8% w/v SDS-PAGE gel and transferred to a Protran BA 83 nylon membrane (0.2  $\mu\text{m}$ , Biorad,  
198 Hercules, USA) with a semi-dry transfer system (Trans-blot SD, Biorad, Hercules, USA). For the  
199 PM H<sup>+</sup>-ATPase blot, a polyclonal antibody against the C-terminal part of the *Arabidopsis*  
200 *thaliana* AHA3 plasma-membrane H<sup>+</sup>-ATPase. The 14-3-3-protein blot was performed using an  
201 antiserum against barley 14-3-3, which was kindly provided by David B. Collinge (Royal  
202 Veterinary and Agricultural University, Frederiksberg C, Denmark). Secondary antibodies (Goat  
203 anti-rabbit IgG alkaline phosphatase conjugate, Biorad, Hercules, USA) were used and the  
204 immunodetection was performed using the standard BCIP/NBT protocol (Promega, Madison,  
205 USA). Relative-intensity band quantifications were determined using ImageJ (1.40g;  
206 <http://rsb.info.nih.gov/ij/>).

207

#### 208 Gene expression analysis

209 At the harvesting times (0, 1, 3, 5, 7, 9 and 11 hours after the initiation of the light period), fully  
210 grown cluster roots were collected, immediately frozen in liquid nitrogen and conserved until  
211 further processing at -80°C. RNA extractions were performed using TRIzol<sup>®</sup> reagent (Invitrogen,  
212 Carlsbad, USA) following manufacturer's instructions and contaminant genomic DNA was  
213 removed with 10 U of DNase I (GE Healthcare, Munich, Germany). Total-RNA samples were  
214 cleaned up using the phenol:chloroform protocol (Maniatis, Sambrook & Fritsch 1989). 1  $\mu\text{g}$  of

215 total RNA (checked for quality and quantity using a spectrophotometer, followed by a migration  
216 in an agarose gel) of each sample was retrotranscribed using 1 pmol of Oligo d(T)<sub>23</sub>VN (New  
217 England Biolabs, Beverly, USA), 15 U Prime RNase Inhibitor (Eppendorf, Hamburg, Germany)  
218 and 10 U M-MuLV RNase H<sup>-</sup> for 1 hour at 42°C (Finnzymes, Helsinki, Finland) following the  
219 manufacturer's instruction. After RNA digestion with 1U RNase A (USB, Cleveland, USA) for 1  
220 hour at 37°C, gene expression analyzes were performed by adding 0.1 µL of the cDNA to  
221 FluoCycle™ sybr green (20 µl final volume; Euroclone, Pero, Italy) in a DNA Engine Opticon  
222 Real-Time PCR Detection (Biorad, Hercules, USA). Primers used (T<sub>m</sub>= 58°C) were the  
223 following: for *LHA1* gene (AY989893), CCATTCATTTCTCTTTTGGGATA and  
224 GAAGACAAAGCTCAATAACCAGAA, for *LHA2* gene (AY989895),  
225 GGAGACTGGCCGAAGACTT and CGGGAATTGAGGCAATACTC, for *LHA3* gene  
226 (AY989894), caggcaattttcceaagaa and acctccagagcaaggcaata, and as housekeeping gene  
227 (polyubiquitin; DQ118117), GCACCCTAGCCGACTACAAC and  
228 CCGGTAAGGGTCTTGACAAA). Triplicates were performed on three independent  
229 experiments; analyses of real-time result were performed using Opticon Monitor 2 software  
230 (Biorad, Hercules, USA) and R (version 2.7.0; <http://www.r-project.org/>) with the qPCR package  
231 (version 1.1-4; <http://www.dr-spiess.de/qpcR.html>). Efficiencies of amplification were calculated  
232 following the authors' indications (Ritz & Spiess 2008): PCR efficiencies were 92.3 and 81.4%  
233 for *LHA1* and *LHA2* genes, respectively. Computation of the graphical representation and  
234 statistical validation (t-test) were performed using SigmaPlot 8.0 (Systat software, Point  
235 Richmond, USA), considering the differences in the PCR efficiency and setting up that *LHA1*  
236 gene expression at time 0 equal 1.  
237

238

## 239 RESULTS

240

### 241 *Citrate, malate and proton exudation by cluster roots*

242 In order to get a better settlement of the diurnal variation of citrate and malate release, we  
243 collected exudates released from fully grown white lupin cluster roots (Fig. 1), corresponding to  
244 the immature and mature stages according to the nomenclature of Massonneau et al. (2001), over  
245 a period of 11 hours after starting the light period (HALS). Figure 2A shows that citrate  
246 exudation progressively increased from 1 HALS to reach a maximum at 5 HALS, where it had  
247 approximately doubled. Thereafter, citrate exudation decreased to levels lower than those  
248 recorded at the beginning of the light period. Exudation rates for malate (Fig 2B) were always  
249 lower than those recorded for citrate and did not change significantly. These results indicate that  
250 there is a differential regulation between citrate and malate exudations.

251 To investigate whether the enhanced citrate exudation was accompanied by changes in root  
252 external medium acidification, fully grown cluster roots were placed on an agar gel containing  
253 the pH indicator bromocresol purple that turns yellow in response to acidification. The time  
254 dependence of agar acidification by cluster roots (Fig. 3) corresponded well with that of citrate  
255 exudation (Fig. 2A). The basal citrate and malate release observed at the beginning or the end of  
256 the light period (Fig. 2, A and B) was not accompanied by detectable acidification of the agar  
257 medium (Fig. 3).

258 In order to verify whether the PM H<sup>+</sup>-ATPase was involved in the development of a higher  
259 citrate exudation rate, cluster roots were treated with fusicoccin, which permanently activates the  
260 plasma-membrane proton pump (Johansson, Sommarin & Larsson 1993; Palmgren 1998).

261 Fusicoccin stimulated proton release from cluster roots (Fig. 3) and was accompanied by an  
262 increase in exudation of citrate, but not of malate (Fig. 2, A and B). The potency of fusicoccin  
263 was strongest in the beginning of the light period and negligible as citrate exudation peaked at 5  
264 HALS (Fig. 2A and 4). These results indicate that activation of the proton pump induces citrate  
265 release, however at 5 HALS, citrate exudation cannot be stimulated further suggesting that the  
266 proton pump does not respond to fusicoccin, *i.e.* it is already operating at maximal activity.

267 Vanadate is a well known inhibitor of P-type ATPases, a family of ion pumps to which the PM  
268 H<sup>+</sup>-ATPase belongs (Cocucci, Ballarin-Denti & Marrè 1980). Application of vanadate to white  
269 lupin cluster roots strongly limited medium acidification (Fig. 3). However, citrate release was  
270 only partially affected by the inhibitor (Fig. 4), the degree of inhibition being dependent on the  
271 time at which cluster roots were treated. At 1 HALS, citrate exudation was almost insensitive to  
272 vanadate (Fig. 4) whereas vanadate was able to completely inhibit the enhanced element of  
273 citrate release, reverting the exudation rate to the level measured at 1 HALS (Fig. 4). This  
274 strongly suggests an involvement of PM H<sup>+</sup>-ATPase in the increased level of citrate release in  
275 cluster roots of P-deficient white lupin.

276

#### 277 *Time-course of cytosolic pH fluctuations*

278 In order to verify whether the burst of citrate and proton release in fully grown cluster roots were  
279 reflected by changes in the cytosolic pH, an *in vivo* NMR spectroscopy approach was used. For  
280 this purpose, the chemical shift ( $\delta$ ) of phosphate (<sup>31</sup>P) resonance was followed. Cytosolic and  
281 vacuolar pH values were calculated using a standard titration curve designed for a “classical”  
282 cellular composition (Roberts, Wadejardetzky & Jardetzky 1981; Spickett, Smirnoff & Ratcliffe

283 1992). Figure 5 illustrates typical NMR profiles measured on roots of P-deficient and P-  
284 sufficient white lupin.

285 Table 1 shows that the cytosolic pH changed paralleling the release of protons and citrate (Fig  
286 2A; Fig. 3). At the beginning of the light period, the cytosolic pH was 7.8 and subsequently  
287 increased to reach a maximum of pH 8.2 after 5 hours where citrate exudation also peaked (Fig.  
288 2A). This alkalization could be a consequence of proton extrusion from the cytosol to the  
289 external solution (Fig. 3). Following the maximum alkalization at 5 HALS, the cytosolic pH  
290 decreased. Addition of fusicoccin similarly resulted in an alkalization of the cytosol (Table 1)  
291 as has also been observed previously (Espen et al. 2000).

292

### 293 *Activity, amount and regulation of the PM H<sup>+</sup>-ATPase*

294 To confirm that the PM H<sup>+</sup>-ATPase was activated in concomitancy with organic acid release,  
295 plasma-membrane vesicles were isolated from fully grown cluster roots at different time points  
296 and characterized their plasma-membrane H<sup>+</sup>-ATPase activities (Fig. 6). Notably, the activity of  
297 PM H<sup>+</sup>-ATPase in vesicles changed during the light period and followed a pattern similar to that  
298 observed for citrate and proton exudation (see Fig. 2A and 3). Thus, it progressively increased up  
299 to 5 HALS and subsequently decreased down to a basal value at 11 HALS. These results clearly  
300 indicate the concomitance of the citrate-exudation burst (Fig. 2A), the acidification of the root  
301 external medium (Fig. 3) and the increase in the ATP hydrolytic activity of the PM H<sup>+</sup>-ATPase  
302 (Fig. 6).

303 In order to determine whether the enhancement of PM H<sup>+</sup>-ATPase activity occurred at the  
304 transcriptional and/or the post-transcriptional level we determined: i) the expression levels of  
305 three PM H<sup>+</sup>-ATPase genes isolated from white lupin roots; ii) the amount of PM H<sup>+</sup>-ATPase

306 protein; iii) the amount of 14-3-3 bound to the plasma membranes; and iv) the pH-dependency of  
307 the PM H<sup>+</sup>-ATPase activity, which is a measure of its regulatory state (Palmgren 1998).

308 Transcriptional activity of PM H<sup>+</sup>-ATPase genes was investigated by real-time RT-PCR analyses  
309 using primers designed from gene sequences of three different H<sup>+</sup>-ATPase isoforms known to be  
310 present in white lupin: *LHA1* (AY989893), *LHA2* (AY989895) and *LHA3* (AY989894). PM H<sup>+</sup>-  
311 ATPase genes showed different expression patterns during the light period (Fig. 7). A distinct  
312 peak in transcript abundance was observed for *LHA1* at 3 HALS; *LHA2* was expressed at a lower  
313 level as compared to *LHA1*, although showing maximal transcript abundance at 7 HALS.  
314 Expression of *LHA3* was not detectable over the whole experimental period (data not shown).

315 In order to investigate whether changes in activity of PM H<sup>+</sup>-ATPase could be explained by  
316 changes in the amount of the corresponding protein, we performed Western blot analyses using  
317 polyclonal antibodies targeted against the C-terminal domain of the enzyme (Fig. 8). Following  
318 the onset of the light period, an increase in the amount of H<sup>+</sup>-ATPase protein was indeed  
319 observed, with a peak at 5 HALS; thereafter, the protein level progressively decreased. The  
320 changes in PM H<sup>+</sup>-ATPase protein levels followed the pattern observed for the ATP hydrolytic  
321 activity of the pump (see Fig. 6).

322 14-3-3 protein is involved in the activation of plasma-membrane proton pump by binding to its  
323 C-terminal domain (Johansson et al. 1993; Palmgren 1998). The possible occurrence of a post-  
324 translational regulation of the plasma-membrane proton pump was analyzed by observing  
325 changes in the amount of activating 14-3-3 protein bound to plasma membranes and by  
326 analyzing the pH-dependency of the enzyme, which moves to a slightly more alkaline maximum  
327 in response to post-translational activation (Palmgren 1998). Indeed, there was a significant  
328 increase in the amount of 14-3-3 protein bound to plasma membranes during the first hours (Fig.

329 8). The pH optimum of PM H<sup>+</sup>-ATPase activity was 6.5 at the beginning of the light period (Fig.  
330 9), which is consistent with the optimum of the down-regulated enzyme (Regenberg et al. 1995).  
331 During the light period, concomitantly with the burst of citrate exudation, the pH optimum  
332 shifted to more alkaline values (6.8 and 7.0 at 3 and 5 HALS, respectively). Taken together,  
333 these results suggest that post-transcriptional regulation of the plasma-membrane proton pump  
334 had occurred at the time of the enhanced release of citrate.

335

336

## 337 DISCUSSION

338

339 Adaptation of white lupin to low-P soils is generally attributed to its capacity to form cluster  
340 roots, from which massive release of carboxylates (citrate and, to a lesser extent, malate) takes  
341 place (Gardner, Parbery & Barber 1981; Keerthisinghe et al. 1998). It has been shown that  
342 carboxylate release occurs according to a developmentally defined program (Massonneau et al.  
343 2001) and may show diurnal variations, with higher rates being recorded during the light period  
344 than in the dark (Watt & Evans 1999).

345 In the present study we have defined in detail the time-course, during the illumination period, of  
346 citrate and malate release rates from immature and mature cluster roots of P-deficient white lupin  
347 (Fig. 1). The results show that the citrate exudation progressively increased from 1 to 5 hours  
348 after the start of the light period (HALS), then decreased to a minimum at 11 HALS (Fig. 2A). A  
349 similar pattern was reported by Hocking and Jeffery (2004) for *Lupinus luteus*. Our data also  
350 show that white lupin cluster roots are able to release malate, but interestingly at lower rates and

351 without showing any significant change during the considered period demonstrating a  
352 differential exudation pattern between citrate and malate (Fig. 2, A and B).

353 A strong acidification of the rhizosphere has been reported to occur concomitantly with the  
354 release of carboxylates (Neumann et al. 2000). Several studies have addressed the question  
355 whether a link exists between carboxylate and proton exudation. Due to the form (anionic) of  
356 carboxylic acids at cytosolic pH values (Ma et al. 2001) and the presence of citrate transport  
357 systems at the plasma membrane of white lupin (Zhang et al. 2004), protons are likely to be  
358 needed for energization and charge compensation during release of carboxylates. Increased  
359 proton release has also been related to the increased cation/anion influx ratio of P-deficient plants  
360 (Shen et al. 2005) and to the presence of the rhizobium-legume symbiosis (Sas et al. 2001).  
361 Activation of the PM H<sup>+</sup>-ATPase has been suggested to be responsible for proton exudation  
362 associated with citrate export in white lupin (Kania et al. 2001; Yan et al. 2002). Recently Zhu et  
363 al. (2005), studying the effect of the PM H<sup>+</sup>-ATPase effectors fusicochin and vanadate on release  
364 of carboxylates (malate and citrate) and several ions (cations and anions) from cluster roots of P-  
365 deficient white lupin, concluded that proton extrusion may serve as charge balance for malate  
366 release, while sodium or potassium might act as counterions for citrate. Results here presented  
367 suggest that the counterions might change in function of how much citrate is released; *i.e.* it  
368 seems there are two different mechanisms of citrate release: one might be an anion channel as  
369 shown by Zhang et al. (2004) and another might be a multidrug and toxic compound extrusion  
370 system (MATE), which transports citrate but not malate, as hypothesized by Furukawa et al.  
371 (2007).

372 In the present study we show that rhizosphere acidification by immature and mature proteoid  
373 roots is concomitant with the burst of citrate exudation (Fig. 3). As expected, proton release was

374 enhanced by fusicoccin treatment. Citrate extrusion could also be enhanced by fusicoccin, but  
375 not when the rate of release was maximal (at 5 HALS, Fig. 2A and 4). This result indicates that  
376 either the proton pump was already fully activated at 5 HALS or that a maximal citrate synthesis  
377 or exudation activity has been reached at that time which could not be further increased by  
378 activation of the PM H<sup>+</sup>-ATPase in response to fusicoccin. After 11 hours of illumination,  
379 fusicoccin had only a negligible effect on citrate release while it was still able to stimulate proton  
380 efflux, suggesting that at this time the internal supply of citrate might have become limiting. On  
381 the other hand, malate release was not affected by the fusicoccin treatment (Fig. 2B). Vanadate  
382 virtually abolished rhizosphere acidification (Fig. 3); on the other hand the effect of the inhibitor  
383 on citrate release was dependent on the time of application, with a decrease of about 50% being  
384 observed at 5 HALS and only of 17% at 1 HALS (Fig. 4).

385 The marked variations in citrate and proton release from the cluster roots during the light period  
386 could conceivably influence the cytosolic pH, therefore we performed time-course measurements  
387 with NMR instrumentation (Fig. 5) and observed that during the burst of citrate and proton  
388 exudation an alkalization of the cytosol took place (Tab. 1), indicating that the cellular pH-stat  
389 could not fully compensate for the release of protons, even if it is known that the capacity to  
390 regulate pH should be increased in white lupin cluster roots; in fact, PEP carboxylase, an  
391 enzyme, which is postulated to play an important role in pH regulation, was shown to be up-  
392 regulated in these tissues (Johnson, Vance & Allan 1996). The lowest pH values were measured  
393 concomitantly with a decrease in proton and citrate efflux (from 7 to 11 HALS). Fusicoccin,  
394 applied at 1 HALS, also led to an increase in cytosolic pH (Tab. 1), as a consequence of PM H<sup>+</sup>-  
395 ATPase activation (Espen et al. 2000).

396 PM H<sup>+</sup>-ATPase activity also changed during the light period (Fig. 6) closely paralleling citrate  
397 and proton extrusion; these modifications were consistent with variations in cytosolic pH. Data  
398 on the cytosolic pH and PM H<sup>+</sup>-ATPase activity further highlight the close link between the  
399 enhanced exudation of proton and citrate in cluster roots of P-deficient white lupin. Modulation  
400 of PM H<sup>+</sup>-ATPase activity have been reported to be due either to altered gene expression or to  
401 post-translational modifications (Michelet et al. 1994; Palmgren 1998; Shen et al. 2005). The  
402 latter type of regulation is achieved by phosphorylation of the C-terminal auto-inhibitory domain  
403 of the PM H<sup>+</sup>-ATPase which enables binding of activating 14-3-3 protein (Fuglsang et al. 1999;  
404 Olsson et al. 1998). In our work changes in PM H<sup>+</sup>-ATPase activity (Fig. 6) strictly concomitant  
405 with the burst of citrate exudation (Fig. 2A) were paralleled by similar changes in protein amount  
406 (Fig. 8); furthermore increased transcript levels for at least one of the genes encoding the three  
407 known enzyme isoforms could be recorded (Fig. 7). These data are consistent with a  
408 transcriptional regulation of the PM H<sup>+</sup>-ATPase. On the other hand, the shift in pH optimum  
409 (Fig. 9) and the pattern of changes in 14-3-3 protein amount (Fig. 8) suggest the occurrence also  
410 of a post-translational regulation of the enzyme. Up-regulation of PM H<sup>+</sup>-ATPase due to  
411 transcriptional and post-translational modifications has been reported for aluminum-stressed  
412 soybean root tips (Shen et al. 2005); furthermore, a transcriptional regulation of PM H<sup>+</sup>-ATPase  
413 involving a specific isoform has also been demonstrated in cucumber plants exposed to Fe-  
414 deficiency (Santi et al., 2005).

415 The present study demonstrates for the first time the involvement of a two-component system in  
416 organic acid exudation in cluster roots of P-deficient white lupin: one, that sustains a basal rate  
417 of malate and citrate release and a second that sustains the burst of citrate exudation which is  
418 closely linked to active proton extrusion from the cytosol. Further, the results indicate that two

419 different mechanisms of organic acid exudation are operating. The first component mediates a  
420 basal rate of malate and citrate release corresponding to about 2  $\mu\text{mol}$  of citrate and 1  $\mu\text{mol}$  of  
421 malate per hour and gram of fresh weight (Fig. 2, A and B, Fig. 4), and is not directly linked to  
422 the PM  $\text{H}^+$ -ATPase activity. The second one corresponds to the burst of citrate release (which is  
423 about two fold higher than the basal citrate release) and is strictly PM  $\text{H}^+$ -ATPase activity  
424 dependent.

425

426

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568

TABLES

**Table I. Variations of cytosolic and vacuolar pH of cluster-root cells during the light period.**

Cluster roots were collected from cluster roots of 5-week-old P-deficient white lupin at different times after the start of the light period. Cytosolic and vacuolar pH values were derived from the chemical shift ( $\delta$ ) of cellular phosphate resonance after construction of a standard titration curve with or without 10  $\mu$ M fusicoccin (FC), using  $^{31}\text{P}$ -NMR in *in vivo* experiments. Values are the means of two independent experiments. Data are presented as differences from the cytosolic pH value determined at 1 HALS which was 7.78. The accuracy is calculated as three times the maximum variation observed in the experiments and is equal to  $\pm 0.02$  pH units for the cytoplasm and  $\pm 0.04$  pH units for the vacuole. nd stand for not detectable.

<b>Hours after light start (HALS)</b>	<b>1</b>	<b>1+FC</b>	<b>3</b>	<b>5</b>	<b>7</b>	<b>9</b>	<b>11</b>
$\Delta$ cytosolic pH	-	0.68	0.13	0.42	-0.07	-0.46	-0.36
$\Delta$ vacuolar pH	-2.51	-2.38	nd	nd	nd	nd	nd

## FIGURE LEGENDS

**Figure 1. Schematic view of a white lupin plant grown for 5-week in a P-deficient nutrient solution and a detailed cluster root.** This figure represent typical cluster root and whole plant of white lupin grown for 5-week in nutrient solution without any external supply of phosphate. Moreover, the different parts of the cluster roots are indicated following the nomenclature of Massonneau et al. (2001). In this work, immature and mature parts were collected together and named fully grown cluster root.

**Figure 2. Citrate (A) and malate (B) secretion from cluster roots during the light period.** Fully grown cluster roots were harvested from 5-week-old P-deficient white lupin at different times after the start of the light period. Exudates were collected after bathing roots for one hour in 10 mM CaSO<sub>4</sub>, 15 mM MES-BTP pH 6 with or without 10 μM fusicoccin. Means ± SD of three independent experiments are reported. Capital letters (A, B) refer to statistically significant differences (t-test, N=3, P<0.05) in organic acid secretion of control roots (- fusicoccin). Asterisks indicate a statistically significant (t-test, N=3, P<0.05) effect of fusicoccin treatment.

**Figure 3. Acidification capacity of cluster roots during the light period.** Fully grown cluster roots were harvested from 5-week-old P-deficient white lupin at different times after the start of the light period. After incubation for 1 hour in 10 mM CaSO<sub>4</sub>, 15 mM MES-BTP pH 6 in the absence or presence of 10 μM fusicoccin or 500 μM vanadate, roots were placed for 30 min on an agar gel containing the pH indicator bromocresol purple and adjusted to pH 6. Yellow and purple colors indicate pH values respectively below and above 6.

**Figure 4. Effect of vanadate and fusicoccin on citrate secretion.** Fully grown cluster roots were harvested from 5-week-old P-deficient white lupin at 1 and 5 hours after the start of the light period (HALS). Exudates were collected after bathing roots for one hour in 10 mM CaSO<sub>4</sub>, 15 mM MES-BTP pH 6 in the absence or presence of 10 μM fusicoccin or 500 μM vanadate. Means + SD of three independent experiments, are reported; relative values in comparison with the untreated control at 1 HALS are also shown. Capital letters (A, B, C) refer to statistically significant differences (t-test, N=3, P<0.05).

**Figure 5. *In vivo* <sup>31</sup>P-NMR spectra of white lupin cluster roots.** Apical segments of lateral roots (the last centimeter) were harvested from plants grown for 5 weeks in the absence (A) or presence (B) of Pi. Each spectrum was recorded using fast acquisition conditions with a recycle time of 1 s and represents the sum of 2,700 scans. The numbered peaks correspond to: 1) cytosolic phosphate; 2) vacuolar phosphate. The region including cytosolic phosphate is shown also on an expanded scale (4x). The chemical shift of the cytosolic phosphate peaks, reflecting the differences in pH values, is highlighted by a vertical dotted line.

**Figure 6. Plasma-membrane H<sup>+</sup>-ATPase activity in cluster roots during the light period.** Plasma-membrane (PM) vesicles were isolated from cluster roots of 5-week-old P-deficient white lupin at different times after the start of the light period. The enzyme activity was measured at pH 6.5. Data are means ± SD of two independent experiments.

**Figure 7. Expression analyses of genes coding for different PM H<sup>+</sup>-ATPase isoforms of cluster roots during the light period.** Total RNA was extracted from cluster roots of 5-week-old P-deficient white lupin at different times after the start of the light period. Relative expression of *LHA1* and *LHA2* genes coding for different PM H<sup>+</sup>-ATPase isoforms of white lupin was analyzed by real-time RT-PCR. Data are means ± SD of three independent experiments. Changes in gene expression were calculated on the basis of expression levels of *LHA1* at 1 hour after the start of the light period (HALS); relative expression of *LHA2* gene was also illustrated in the insert with a higher magnification. Asterisks indicate a statistically significant (t-test, N=3, P<0.05) difference in expression level of each gene with respect to its expression at 1 HALS.

**Figure 8. Western-blot analysis of the PM H<sup>+</sup>-ATPase and 14-3-3 proteins in cluster roots during the light period.** Purified plasma membranes isolated from cluster roots of 5-week-old P-deficient white lupin at different times after the start of the light period were used for immunodetection of PM H<sup>+</sup>-ATPase using antibodies raised against the C-terminal part of the *Arabidopsis* AHA3 PM H<sup>+</sup>-ATPase and of 14-3-3-protein using antibodies raised against the barley 14-3-3 protein. Quantification of western-blot signals is expressed as percentage of that recorded at 1 hour after the start of the light period (HALS). Data of a representative experiment are reported.

**Figure 9. pH dependency of PM H<sup>+</sup>-ATPase activity of cluster roots during the light period.** Plasma-membrane (PM) vesicles were isolated from cluster roots of 5-week-old P-deficient white lupin at 1, 3, 5, 9 hours after the start of the light period (HALS). The enzyme activity was measured at different pH values in the range between 6.2 and 8.0. Data are means ± SD of two independent experiments.

Figure 1.

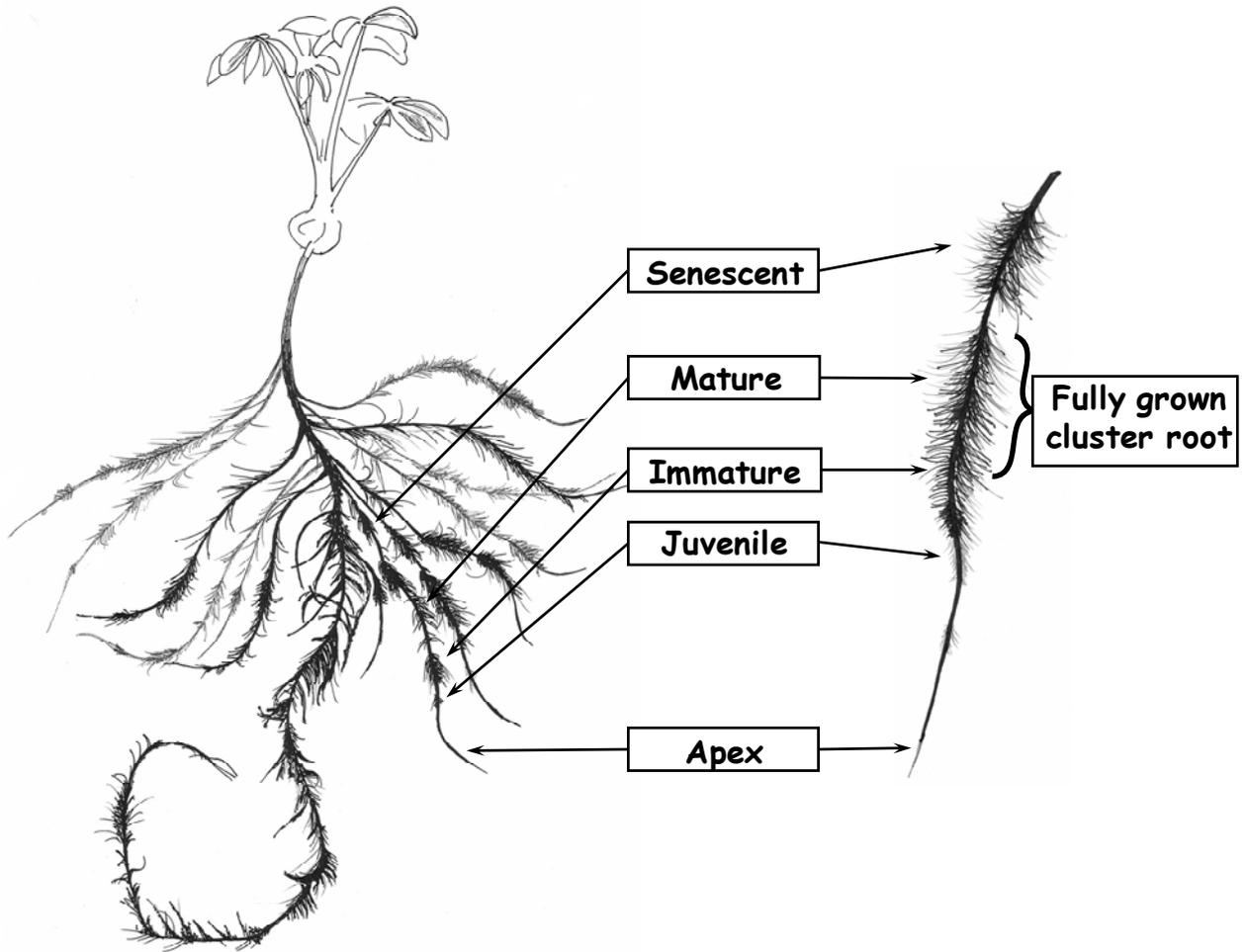


Figure 2.

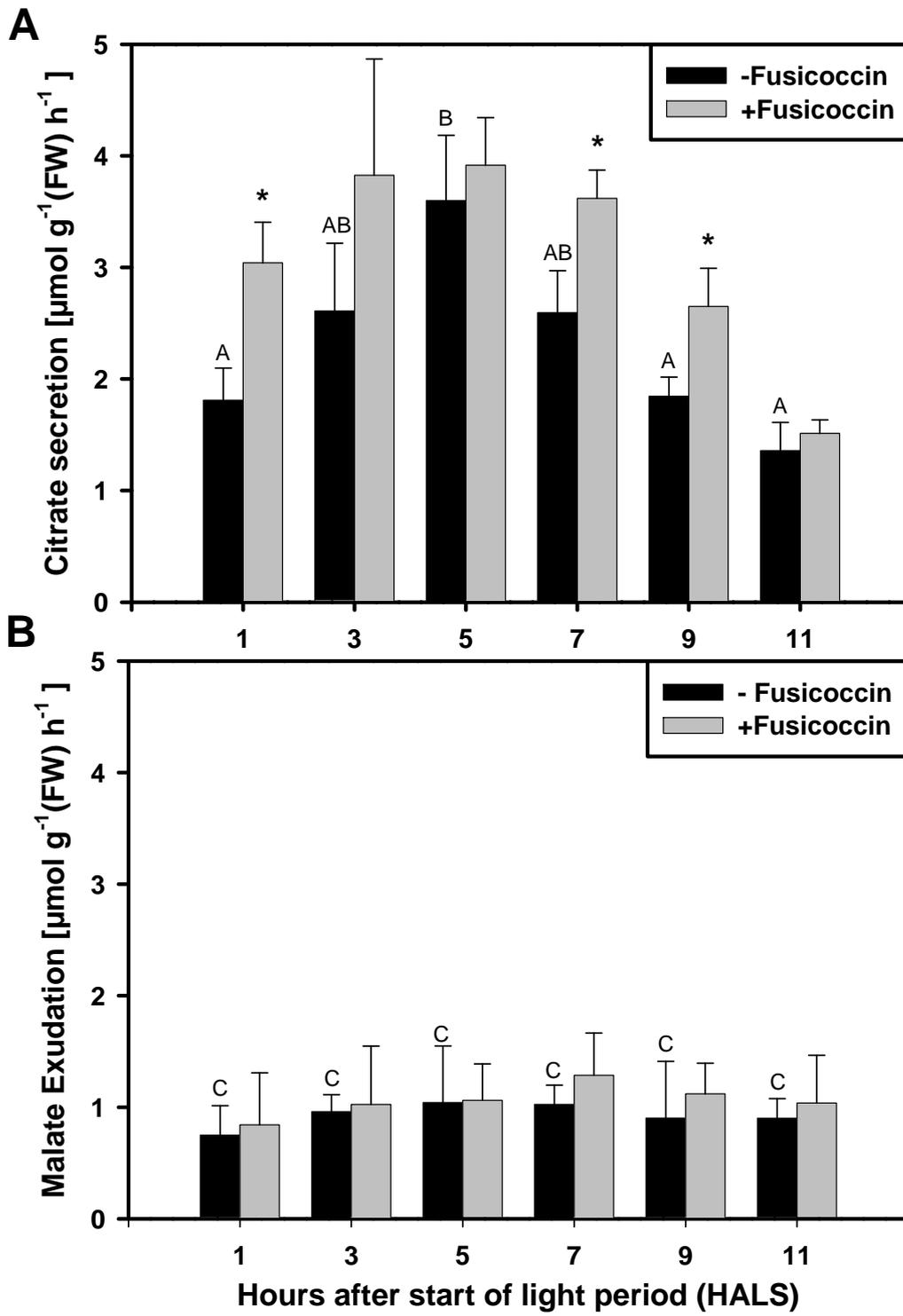


Figure 3.

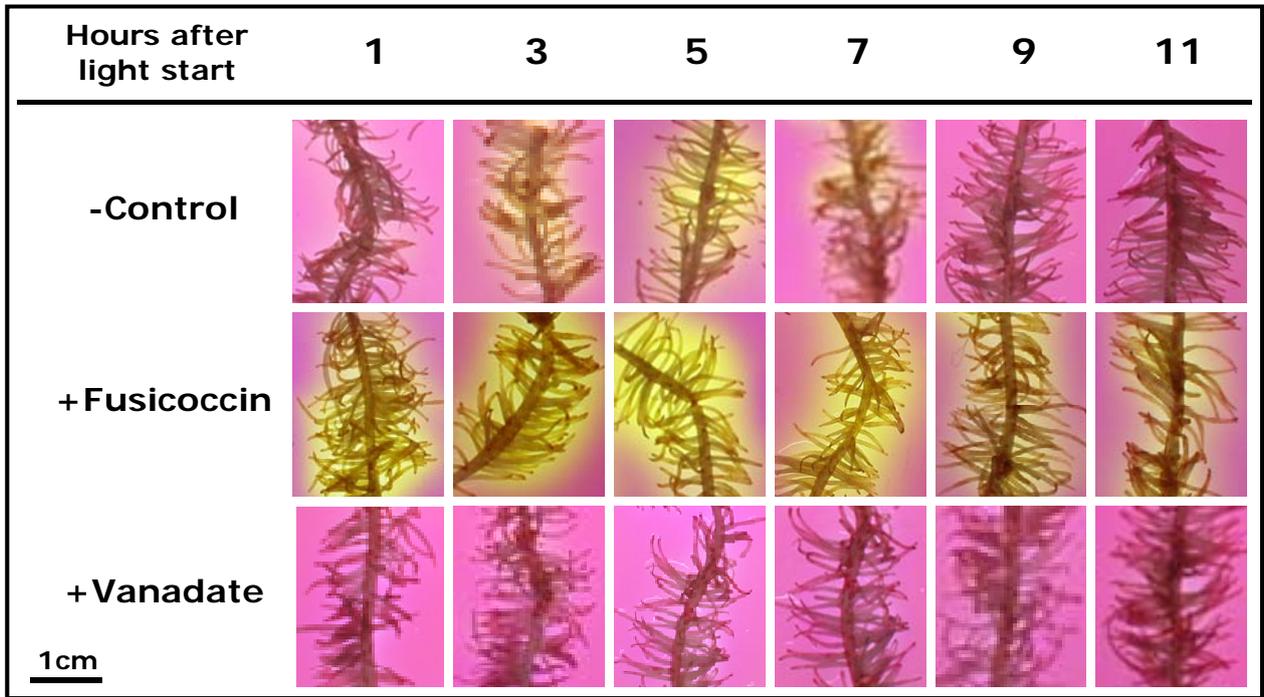


Figure 4.

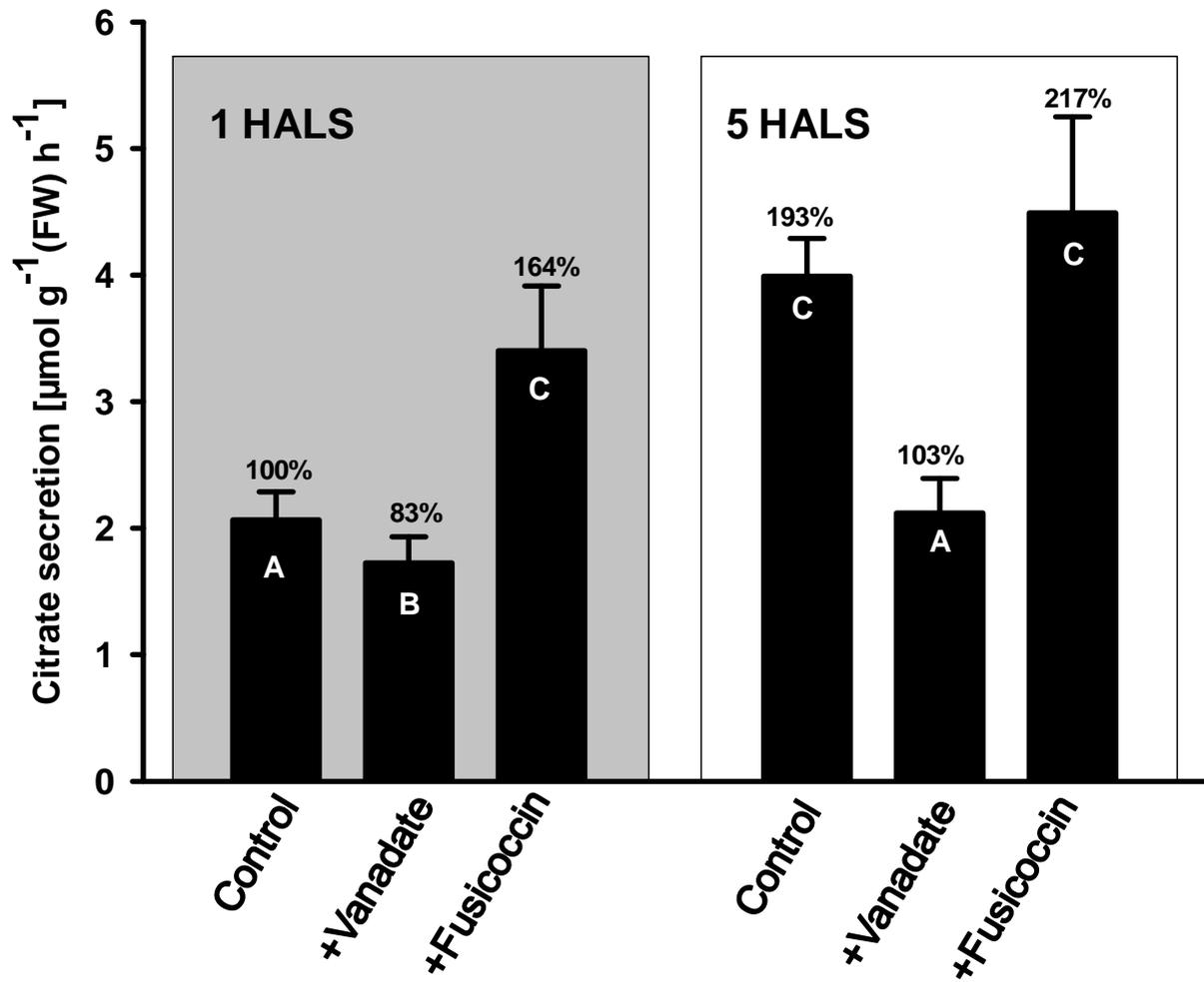


Figure 5.

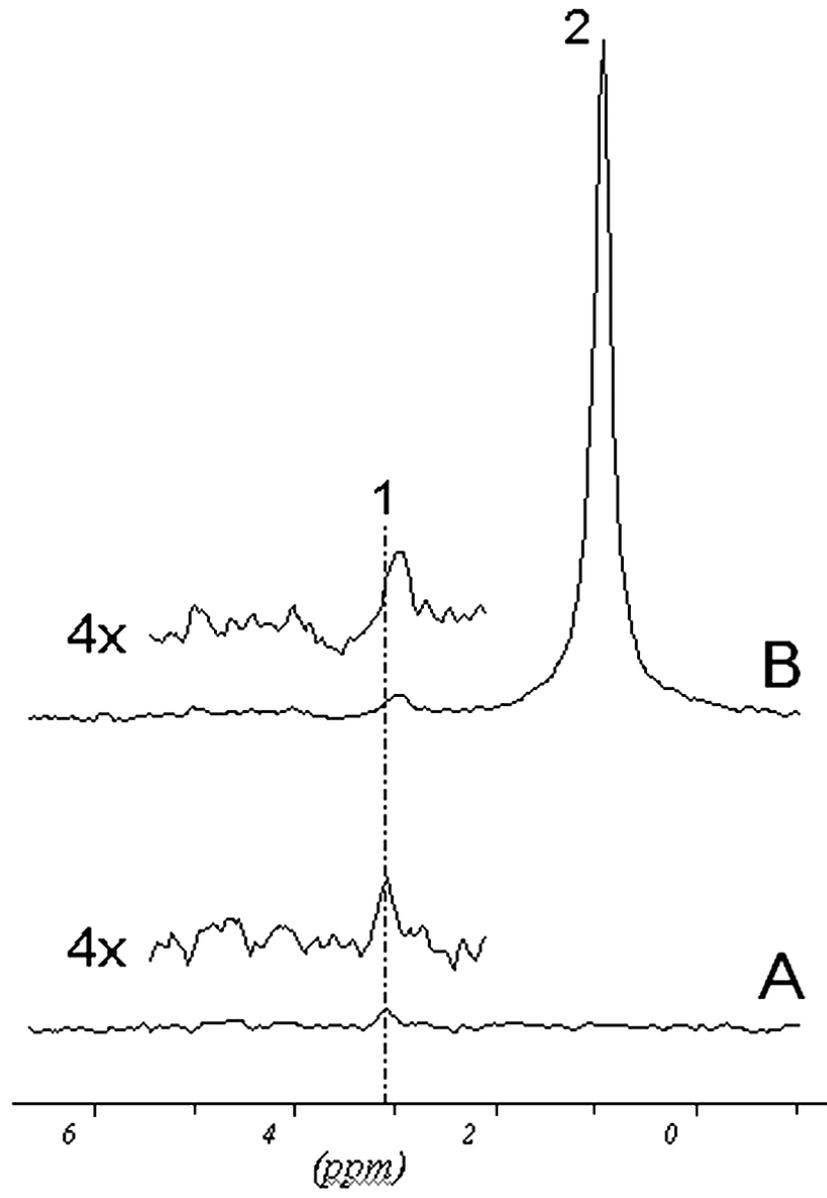


Figure 6.

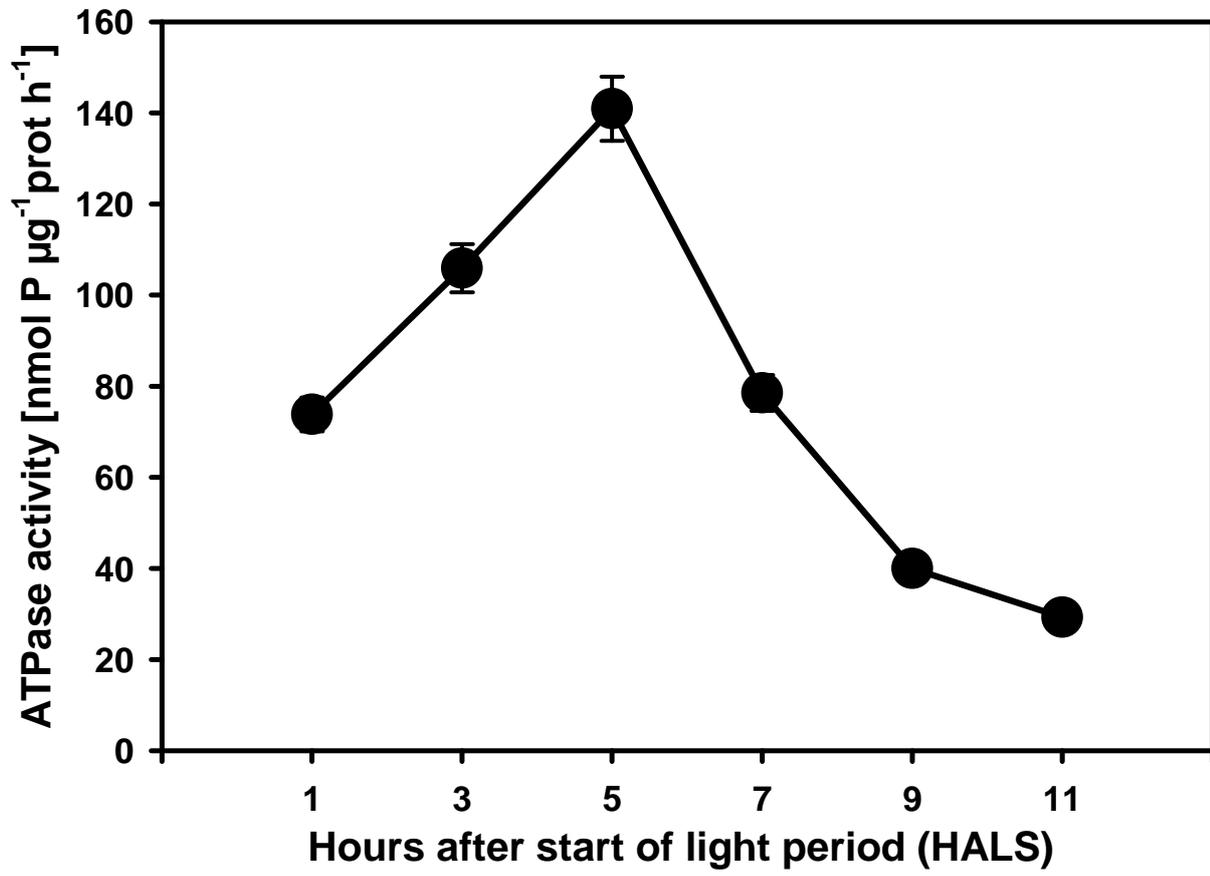
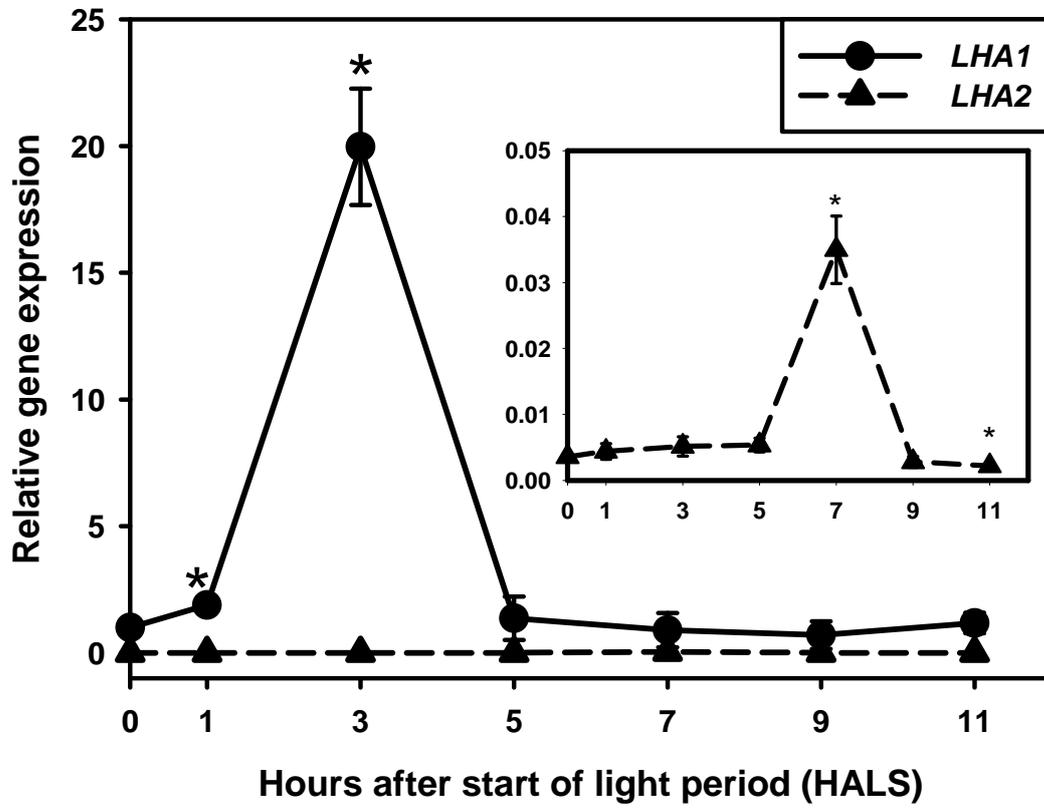


Figure 7.



**Figure 8.**

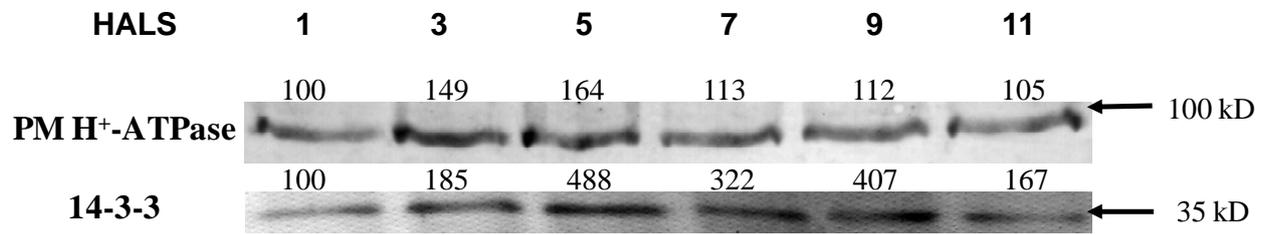
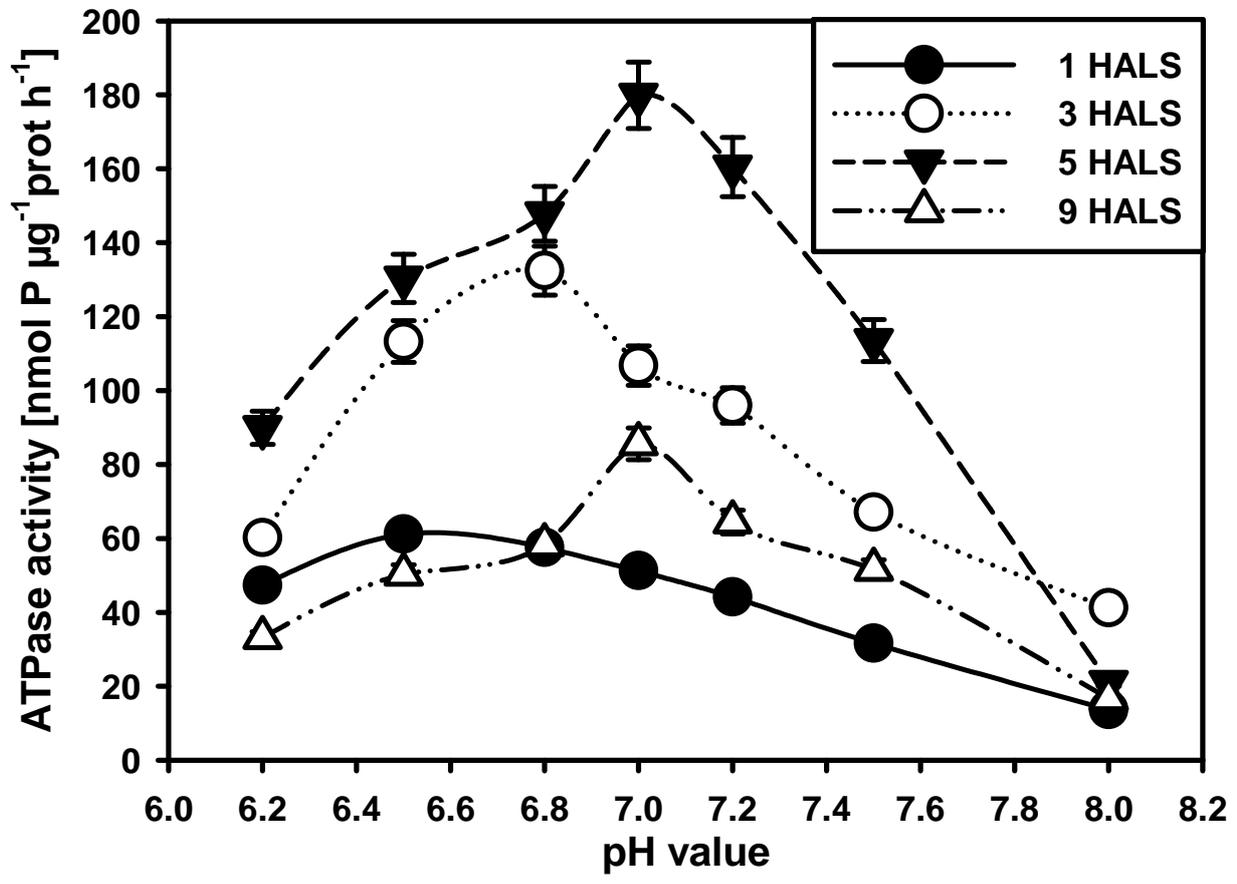


Figure 9.





**Effects of PM  $H^+$ -ATPase modulators on proton extrusion by cluster roots at the time of maximum exudation.** Fully grown cluster roots were harvested from 5-week-old P-deficient white lupin at 5 hours after the start of the light period. After incubation for 1 hour in 10 mM  $CaSO_4$ , 15 mM MES-BTP (pH 6) in the absence or presence of 10  $\mu$ M fusicoccin or 500  $\mu$ M vanadate, roots were placed for 30 min on an agar gel containing the pH indicator bromocresol purple and adjusted to pH 6. Yellow color indicates pH values below 6.