RESEARCH PAPER

2	Intra-cellular distribution and binding state of aluminium in root apices of two common
3	bean (Phaseolus vulgaris) genotypes in relation to Al toxicity
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12	Running title: Intra-cellular distribution of Al and its relationship to Al toxicity
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1 Abstract

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3 The role of the intra-cellular distribution and binding state of aluminium (Al) in Al toxicity, 4 using Al exchange and Al fractionation methodologies, were studied in two common bean 5 (Phaseolus vulgaris L.) genotypes differing in Al resistance. These two genotypes are 6 characterized by a similar initial period (4 h) of Al sensitivity followed by a contrasting 7 recovery period (8-24 h). A higher initial Al accumulation in Quimbaya (Al-resistant) in the 5 8 mm root apex compared to VAX-1 (Al-sensitive) could be related to its higher content of 9 unmethylated pectin and thus higher negative charge of the cell walls. The binding state and 10 cellular distribution of Al in the root apices revealed that the root elongation-rate was 11 significantly negatively correlated with the free apoplastic and the stable-bound, not citrate-12 exchangeable cell-wall Al representing the most important Al fraction in the root apex (80%), 13 but not with the symplastic and the labile-bound, citrate-exchangeable cell-wall Al. It is 14 postulated that the induced and sustained recovery from the initial Al stress in the Al-resistant 15 genotype Quimbaya requires reducing the stable-bound Al in the apoplast thus allowing cell 16 elongation and division to resume.

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18 **Key words**: aluminum toxicity, apoplast, cell wall, compartmentation, root apex

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1 Abreviations

2	CW	Cell wall
3	CEZ	Central elongation zone
4	DM	Degree of methylation
5	DTZ	Distal transition zone
6	EZ	Elongation zone
7	Al _{mono}	Monomeric aluminium
8	PEM	Pectin methylesterase
9	PCV	Pyrocatechol violet
10	SYM	Symplast
11	ΤZ	Transition zone
12	WFSF	Water free space fluid
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1 Introduction

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3 Aluminium (Al) toxicity is a major factor limiting plant growth especially on acid soils in the 4 tropics and subtropics (von Uexküll and Mutert 1995). Common bean (Phaseolus vulgaris L.) 5 growing areas of about 40% of Latin America and 30 to 50% of central, eastern, and southern 6 Africa are affected by Al toxicity resulting in yield reduction from 30 to 60% (CIAT 1992). The primary effect of Al is an inhibition of root growth (Foy 1988), an effect that can be seen 7 8 within hours of Al treatment (Llugany et al. 1995, Blamey et al. 2004). The major site of Al 9 perception and response is the root apex (Ryan et al. 1993), and particularly the distal part of 10 the transition zone (DTZ, 1-2 mm) is the most Al-sensitive apical root zone (Sivaguru and 11 Horst 1998, Kollmeier et al. 2000). In common bean in contrast to maize (Zea mays L.), 12 however, Al applied to the elongation zone (EZ) contributed to the overall inhibition of root 13 elongation by Al (Rangel et al. 2007). Also, common bean differs from most other plant 14 species, particularly cereals, through a lag phase after the beginning of Al treatment before Al 15 resistance mechanisms are expressed (Cumming et al. 1992, Rangel et al. 2007). This is 16 typical for a pattern II response to Al treatment (Ma et al. 2001) characterized by an Al-17 induced delayed (several hours) exudation of organic acid anions, particularly citrate in 18 common bean (Mugai et al. 2000, Ma et al. 2001, Shen et al. 2002, Rangel and Horst 2006, 19 Stass et al. 2007).

The role of the root exudation of organic acid anions in reducing Al uptake/binding in the root apoplast thus enhancing Al resistance is widely accepted particularly in pattern I plant species (Ma et al. 2001, Ryan et al. 2001, Kochian et al. 2004, Delhaize et al. 2007). However, the role of symplastic lesions of Al toxicity and of sequestration of Al by organic ligands as a mechanism of Al resistance are still issues of debate (Vázquez et al. 1999, Illes et al. 2006). Thus, there is a need to better understand the kinetics of Al accumulation in root apices and its distribution at a cellular and tissue level in relation to genotypic differences in Al resistance
 particularly in pattern II plant species such as common bean.

3 Aluminium accumulates in roots with a rapid initial phase (accumulation of easily 4 exchangeable Al in the apoplast) followed by a lower linear rate (metabolism-dependent 5 binding of Al into the apoplast and transport of Al into the symplast, Zhang and Taylor 1989, 6 1990). In the apoplast, the negative charge of the cell wall (CW) established by the pectin 7 content and its degree of methylation is a major determinant of this initial Al accumulation 8 (Blamey et al. 1990, Grauer and Horst 1992, Schmohl and Horst 2000, Schmohl et al. 2000) 9 and Al injury (Schmohl et al, 2000, Eticha et al. 2005a, Horst et al. 2007) through altering 10 CW characteristics and functions, such as extensibility, porosity, hydraulic conductivity, 11 displacement of ions from critical sites (Rengel 1990, Blamey et al. 1993, Mimmo et al. 2003, 12 Sivaguru et al. 2006, Horst et al. 2007) and/or disrupting the CW-plasma membrane-13 cytoskeleton continuum (Sivaguru et al. 1999, Horst et al. 1999).

14 There is no doubt that Al can enter the symplast (Tice et al. 1992, Lazof et al. 1994, Vázquez 15 et al. 1999, Eticha et al. 2005b). Taylor et al. (2000) using the model giant algae Chara 16 corallina showed that Al can be transferred from the apoplast to the symplast. However, the 17 low rates of transport observed through the plasma membrane will favor the accumulation of 18 Al in the apoplast (Rengel and Reid 1997). Therefore, interactions of Al with the CW and 19 plasma membrane will necessarily precede any transport into the symplast, these interactions 20 being potentially harmful (see above, Delhaize and Ryan 1995). According to the above 21 scenario, internalization of Al in the symplast (Vázquez et al. 1999, Illes et al. 2006) appears 22 to be a mechanism of Al resistance rather than of Al toxicity.

The main objective of the study was to elucidate the role of the intra-cellular distribution and binding state of aluminium (Al) in relation to Al toxicity in two common bean genotypes differing in Al resistance. These two genotypes appeared to be particularly suitable for this study because they are characterized by a similar initial period of Al sensitivity followed by a
 contrasting recovery period (Rangel et al. 2007).

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4 Materials and methods

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6 Plant material and growth conditions

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8 Seeds of the Al-resistant common bean genotype Quimbaya, large seeded Andean cultivar, 9 and an Al-sensitive genotype VAX-1, small seeded Mesoamerican advanced line (Rangel et 10 al., 2005) kindly supplied by the Bean Outcome Line of CIAT (International Center for 11 Tropical Agriculture, Cali, Colombia) were germinated between filter-paper and foam 12 sandwiches soaked with tap water in an upright position. Uniform seedlings were transferred 13 to 181 pots with constantly aerated simplified nutrient solution (Rangel et al. 2005). Plants 14 were cultured in a growth chamber with controlled environmental conditions of a 16/8 h 15 light/dark regime, 27/25°C day/night temperature, 70% relative air humidity, and a photon flux density of 230 μ mol m⁻² s⁻¹ photosynthetic active radiation at the plant level (Sylvania 16 17 Cool White, 195 W, Philips, Germany).

After 24 h the pH of the solution was lowered gradually from 5.6 to 4.5 and kept constant throughout the treatment period using an automatic pH titration device with 0.1 M HCl/KOH. Plants were treated with 0 or 20 μ M AlCl₃ for up to 24 h. Mononuclear Al (Al_{mono}) concentrations were measured colorimetrically using the pyrocatechol violet method (PCV) according to Kerven et al. (1989). Nominal 20 μ M Al supply resulted in 16 ± 2 μ M Al_{mono} after 24 h. Then the roots were washed in distilled water and the root tips (5 mm length) were harvested for pectin and Al determinations.

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26 Effect of Al on root growth

Two hours before Al treatment tap roots were marked 3 cm behind the root tip using a fine point permanent marker (Sharpie blue, Stanford) which did not affect root growth during the experimental period. Afterwards, the plants were transferred to simplified nutrient solution (Rangel et al. 2005) containing 0 or 20 μ M AlCl₃. Root elongation was measured at 4, 8 and Al treatment using a 1 mm scale.

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8 Determination of pectin and its degree of methylation

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10 Pectin and its degree of methylation was determined in 25 (5 mm) root tips per sample, which 11 were excised and collected in 96% (v/v) ethanol in Eppendorf vials. Root samples were 12 thoroughly homogenized in ethanol using a mixer mill (MM200; Retsch, Haan, Germany) at a 13 speed of 30 oscillations per second for 3 min. The homogenization was repeated twice. Cell-14 wall material was prepared as alcohol-insoluble residue after repeated washing with ethanol, 15 modified after Schmohl and Horst (2000). After every ethanol addition, the samples were 16 centrifuged at 23000 g for 10 min and the supernatant was discarded. The remaining CW was 17 dried using a centrifugal evaporator (RC10-22T, Jouan SA, France), weighed, and hydrolysed 18 according to Ahmed and Labavitch (1977) extending the incubation time to 10 min in 19 concentrated H₂SO₄ and the hydrolysis completed overnight by a stepwise dilution with 20 double-deionized water. The uronic acid content was determined colorimetrically according to 21 Blumenkratz and Asboe-Hansen (1973) using a microplate spectrophotometer (μ QuantTM, 22 Bio-Tek Instruments, Winooski, Vermont, USA). Galacturonic acid was used as a calibration 23 standard, thus the root pectin content is expressed as galacturonic acid equivalents (GalA). 24 For the determination of the degree of methylation (DM), the CW was prepared in the same 25 way as for pectin determination. Methanol was released from the CW by saponification

26 according to Fry (1988), modified after Wojciechowski and Fall (1996). After addition of 2

1	units of alcohol oxidase (EC 1.1.3.13 from Piccia pastoris Sigma, Deisenhofen, Germany) the
2	complex of formaldehyde with Fluoral-P (15 mg ml ⁻¹) (Molecular Probes, Leiden, The
3	Netherlands) was measured fluorometrically (excitation $\lambda = 405$ nm, emission $\lambda = 503$ nm).
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5	Aluminium exchange from intact root tips

7 For the exchange (desorption) of Al from the root tips, roots of twelve seedlings were quickly 8 washed with double-deionized water, then 5 mm root tips were excised with a razor blade and placed in filter units with a pore size of 0.45 µm (GHP Nanosep[®] MF Centrifugal Device, Pall 9 10 Life Sciences, Ann Arbor, USA). Loosely bound Al was exchanged with 500 µl of 50 mM 11 BaCl₂, for 15 min. Root tips were briefly washed in 500 µl of double-deionized water and 12 then transferred for 15 min to 500 μ l of 33 mM Na₃-citrate (pH 5.8) and the filtrate collected 13 in a new vial. Preliminary experiments had shown that longer incubation periods did not 14 release more Al in either fraction. Desorption experiments were conducted at 4°C to minimize 15 loss of Al from the symplast (Zhang and Taylor 1989). Thereafter, the root tips were washed 16 and transferred into a new Eppendorf vial for Al determination.

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18 Aluminium fractionation

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For the determination of apoplastic and symplastic Al fractions in the root tips, the apoplastic and symplastic saps from root tips were collected according to the method described by Yu et al. (1999) and modified by Wang et al. (2004). Briefly, freshly excised 5 mm root tips from twenty five seedlings were arranged in a filter unit (Ultrafree-MC, 0.45 μm; Millipore, Bedford, MA) with the cut ends facing down, and the water free-space fluid (WFSF) collected by centrifugation (4000 g) at 4°C for 15 min. After collecting the WFSF, the root tips were frozen at -20°C. The first symplastic-Al fraction (SYM-1) was recovered from the frozen-

1 thawed samples by centrifugation (4000 g) at 4°C for 15 min. The residue was then 2 transferred to Eppendorf vials and homogenized in 500 µl of ethanol with a mixer mill 3 (MM200; Retsch, Haan, Germany) at a speed of 30 oscillations per second for 3 min. All 4 further centrifugation steps were conducted at 23000 g (4°C) for 5 min. After centrifugation, 5 supernatant and pellet were separated and the pellet suspended again in 500 µl of ethanol. The 6 complete process was repeated twice and both supernatants combined. The supernatants 7 representing the second symplastic-Al fraction (SYM-2) were evaporated in a centrifugal 8 evaporator (RCT 10-22T; Jouan, Saint-Herblain, France) for later Al determination. 9 Subsequently, the pellet consisting of the CW was desorbed at room temperature with 500 µl 10 of 33 mM Na₃citrate (pH 5.8) for 15 min. After centrifugation, the supernatant containing the 11 labile-bound CW Al fraction was analysed for Al. The pellet was washed with double-12 deionized water, centrifuged and the supernatant discarded. Thereafter, the pellet containing 13 the stable-bound CW Al fraction was dried in a centrifugal evaporator (RCT 10-22T; Jouan, 14 Saint-Herblain, France) for later determination of Al.

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16 Determination of Al

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For Al determination, roots, CW material, and the SYM-2 fractions were digested in 500 μ l ultra-pure HNO₃ (65%) overnight. Digestion was completed by incubation in a water bath at 80°C for 20 min. BaCl₂-exchangeable, citrate-exchangeable, WSWF, and SYM-1 Al fractions were directly measured using a Unicam 939 QZ graphite furnace atomic absorption spectrophotometer (GFAAS; Analytical Technologies Inc., Cambridge, UK) at a wavelength of 308.2 nm and with an injection volume of 20 μ L. When required, the samples were diluted with double-deionized water.

1 Statistical analysis

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3 Each experiment had a completely randomized design with four replicates. The ANOVA 4 procedure of the statistical program SAS 9.1 (SAS Institute, Cary, NC, USA) was used for 5 analysis of variance. Means were compared using the Tukey test. *, **, *** denotes 6 significant differences at p < 0.05, 0.01, and 0.001, respectively; n.s. denotes not significant. 7 8 Results 9 10 Effect of Al on overall root elongation 11 12 In presence of Al root elongation of both genotypes was severely inhibited (60-65%) 4 h after 13 the beginning of the Al treatment (Fig. 1a). After 8 h Al treatment, both genotypes recovered, 14 Quimbaya more than VAX-1. Whereas this recovery continued in Quimbaya until the root-15 elongation rate nearly reached the level of the control (without Al), VAX-1 was increasingly 16 damaged by Al after 24 h of Al treatment which is reflected by the highly significant 17 genotype x time interaction. 18 19 Effect of Al treatment on Al content in root tips 20 21 The decrease in root-elongation rate after 4 h Al supply in both genotypes was associated with 22 an increase in Al content in the root tips (Fig. 1b). Recovery of root-elongation rates after 8 h 23 of Al treatment was accompanied by reduced Al contents in the root tips. This decrease 24 continued during further recovery in the Al resistant genotype (Quimbaya), while Al contents 25 increased again in the Al-sensitive genotype (VAX-1) after 24 h of Al treatment (highly

significant genotype x time interaction). Aluminium contents per unit root tip length (5 mm)

after 4 and 8 h of Al treatment were significantly higher in Quimbaya than in VAX-1 (86%). When the Al contents were expressed on a root tip fresh weight basis (nmol per mg root tip; data not shown), this difference was somewhat lower (about 70%) due to a higher mass of the root tips of Quimbaya (9.55 \pm 1.1 mg) as compared with VAX-1 (8.70 \pm 2.4 mg).

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6 Fig. 1

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8 Determination of pectin and its degree of methylation

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10 Since binding of Al to CW is mainly due to pectins, the pectin content and its degree of 11 methylation was determined in the CW isolated from 5 mm root tips. Constitutively, 12 Quimbaya had significantly higher CW pectin contents than VAX-1 (Fig. 2a) whereas the 13 DM did not differ between the two genotypes before the start of the Al treatment (Fig 2b). 14 Aluminium treatment increased the pectin contents in both genotypes independent of Al 15 treatment duration (Fig. 2a) while the DM decreased after 4 h Al treatment in both genotypes 16 (Fig. 2b). However, while recovery of root growth in Quimbaya at longer Al-treatment 17 duration was reflected by increased DM up to the initial value, it remained at the lower level 18 in VAX-1. The resulting content of unmethylated pectin (Fig. 2c) which is a measure of the 19 negativity of the CW was consistently higher (31%) in Quimbaya than in VAX-1. This 20 genotypic difference was smaller (17%) but still significant when the pectin contents were 21 expressed on a CW mass basis (nmol per mg CW, data not shown) due to a higher mass of 22 CW recovered from the twenty five root tips of Quimbaya (4.5 \pm 0.2 mg) as compared to 23 VAX-1 (3.6 ± 0.1 mg). After 24 h Al treatment which did not affect the dry mass of CW per 24 root tip, the content of unmethylated pectin decreased again corresponding to the observed 25 recovery in DM to the level observed prior to the Al treatment in Quimbaya but not in VAX-

This is reflected by the significant genotype x time interaction observed in DM and
 unmethylated pectin.

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4 Fig. 2

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6 Binding state of Al in the root tips

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8 The recovery from initial Al stress leading to genotypic differences in Al resistance might be 9 related to changes and differences in the binding state and compartmentation of Al in the root 10 apices. Therefore, in a first approach root apices were subjected to a fractionated desorption 11 of Al in order to differentiate between loosely and firmly bound Al. Excised root tips were 12 incubated for 15 min each in 50 mM BaCl₂ and then in 33 mM Na₃citrate in order to release 13 from the apoplast free and exchangeable bound Al, and Al weakly bound to the unmethylated 14 pectin, respectively. The Al that was not released from the root tips was considered as non-15 exchangeable (symplastic and more strongly bound apoplastic Al). BaCl₂ was not able to 16 release any detectable amounts of Al from the root tips (data not shown). Incubation in 17 Na₃citrate released between 10 and 30% of the total Al in the root tips depending on the 18 genotype and the Al-treatment period (highly significant genotype x time interaction, Fig. 3). 19 In both genotypes the decrease of root elongation after 4 h Al treatment (see Fig. 1a) was 20 characterized by substantial Al accumulation in both Al fractions (Fig. 3). During the 21 recovery from Al injury after 8 h, Al contents decreased more in the non-exchangeable than in 22 the Na₃citrate-exchangeable fraction. The Al contents of both fractions were higher in 23 Quimbaya than in VAX-1 during the first 8 h. After 24 h Al treatment the picture changed. 24 Whereas in Quimbaya the Al contents continued to decrease in both fractions, they increased again only in the non-exchangeable fraction in VAX-1, leading to a highly significant 25 26 genotype x time interaction.

The total Al contents in the root tips were only loosely related to the root elongation rates when calculated across genotypes and Al treatment duration ($r^2 = 0.22^*$). However, in Quimbaya the increase in root elongation during the recovery from initial Al stress was highly significantly related to both Na₃citrate-exchangeable and non-exchangeable Al (Fig. 4). This was also true for VAX-1 for the recovery period of 4-8 h after Al treatment. However, the severe inhibition in root elongation after 24 h Al treatment in VAX-1 appears to be mainly due to an increase in non-exchangeable Al rather than Na₃citrate-exchangeable Al.

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9 Fig. 3

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11 Fig. 4

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13 In order to better characterize the intra-cellular distribution of Al in the root apices in addition 14 to the binding state, the root tips were subjected to a more differentiated fractionation 15 procedure (Fig. 5). The Al contents in the WFSF fraction were initially two times higher in 16 Quimbaya than in VAX-1 which was due equally to both a greater WFSF volume (estimated 17 on the basis of the recovered amount of WFSF by centrifugation) and Al concentration. The 18 WFSF Al fraction decreased over time in Quimbaya because of decreasing Al concentration 19 rather than volume, whereas the level remained constant in VAX-1, leading to comparable Al 20 contents in both genotypes after 24 h Al treatment (highly significant genotype x time 21 interaction). The SYM-1 Al fraction had a similar order of magnitude as the WFSF Al 22 fraction. Again, the Al contents were higher in Quimbaya than in VAX-1. This fraction did 23 not change with the Al treatment duration in either genotype. The SYM-2 Al fraction which 24 was about twice as high as the two previous fractions was significantly higher after 24 h Al supply to Quimbaya, too. It increased up to 8 h after Al treatment and then remained at this 25 26 higher level in both genotypes. The only fraction which was consistently higher in the Alsensitive VAX-1 was the labile-bound CW Al fraction. This fraction remained stable in Quimbaya but steadily increased in VAX-1 up to 24 h Al treatment. The stable-bound CW Al fraction was quantitatively the most important Al fraction. Initially (up to 8 h Al treatment) the Al contents were higher in Quimbaya but readily decreased with time, whereas in VAX-1 the contents drastically increased after 24 h of Al treatment.

6 Fig. 5

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8 The stable-bound CW Al fraction represented about 80% of the total Al content in both 9 genotypes (Fig. 6). Whereas the relative contribution of this fraction to the total Al content 10 decreased in Quimbaya with time, in VAX-1 this fraction first decreased (8 h) but then 11 increased again after 24 h Al treatment. The second greatest Al fraction was the symplastic 12 fraction (combining the two symplastic Al fractions). This fraction became increasingly 13 greater with Al treatment duration in Quimbaya, while in VAX-1 this was only the case up to 14 8 h. Later, this fraction decreased again. The relative contribution of the WFSF (smallest 15 fraction) and the labile-bound CW Al fractions did not vary much over time. However, the 16 latter was greater in VAX-1 (8%) than in Quimbaya (4.5%).

17 As in the Al-exchange experiments (see Fig. 4), the correlations between root-elongation rate 18 and apoplastic (WFSF and CW Al fractions combined) and symplastic Al contents were 19 calculated separately for each genotype because of the highly significant genotype x time 20 interaction for most Al fractions (see Fig. 5). In both genotypes, root-elongation rate was 21 negatively related to the Al content of the apoplast (Fig. 7a). This is clearer in Quimbaya with 22 a continuous recovery from first Al injury at 4 h than in VAX-1, where the initial recovery 23 after 8 h is followed by severe Al injury after 24 h Al treatment. Symplastic Al was not 24 related to root elongation rate in either genotype. There was even a tendency of a positive 25 correlation.

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- 3 Fig. 7
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5 In order to clarify whether the recovery of root elongation from initial Al stress in Quimbaya 6 can be related to changes in specific apoplastic Al fractions, correlation coefficients were 7 calculated (Fig. 8). The WFSF and the stable-bound CW, but not the labile-bound CW Al 8 fractions showed a highly-significant negative relationship with the enhanced root-elongation 9 rate during the Al treatment.

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11 Fig. 8

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13 **Discussion**

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The results from this study clearly provide additional physiological and biochemical evidence needed to substantiate the previous finding that Al resistance in common bean is an Alinducible trait involving a lag phase of 4-6 hours in contrast to many other plant species (Cumming et al. 1992, Rangel et al. 2007; Fig. 1a).

19 Aluminium treatment resulted in rapid Al accumulation (after 4 h, Fig. 1b) more in the Al-20 resistant genotype Quimbaya than in the Al-sensitive genotype VAX-1 leading to severe 21 decrease in the root-elongation rates in both genotypes (Fig. 1a). Aluminium is accumulated 22 by roots with a rapid initial phase and a lower rate, thereafter (Zhang and Taylor 1989, 1990). 23 The primary binding site of Al is likely the pectic matrix of the CW with its negatively 24 charged carboxylic groups having a particularly high affinity for Al (Blamey et al. 1990, 25 Chang et al. 1999). Short-term Al accumulation by roots is closely related to the pectin 26 content. This may explain the differences in initial Al accumulation between monocots and

1 dicots (Schmohl and Horst 2000, Horst et al. 2007) with their higher CW pectin content 2 (Carpita and Gibeaut 1993). In fact, the factor responsible for Al binding to pectin is not the pectin content but its negative charge determined by its DM which is controlled by pectin 3 4 methylesterase (PME) (Bordenave 1996, Gerendás 2007). The role of the CW pectin-content 5 and its DM in Al resistance has been demonstrated in maize (Schmohl et al. 2000, Eticha et 6 al. 2005a), potato (Solanum tuberosum L., Schmohl et al. 2000, Horst et al. 2007) and 7 common bean (Stass et al. 2007) using different experimental approaches. Eticha et al. 8 (2005*a*) showed that the higher Al accumulation into the root apex in the maize cultivar Lixis 9 was related to its higher content of low-methylated pectin and thus higher negativity of the 10 cell wall compared to the cultivar ATP-Y. Therefore, it appears reasonable to assume that the 11 higher Al accumulation of the common bean genotype Quimbaya in the root apex compared 12 to VAX-1 (Fig. 1) is due to its higher content of unmethylated pectin (higher negative charge 13 of the CW, Fig. 2).

However, in contrast to maize where a higher negative charge of the CW contributes to genotypic Al sensitivity (Eticha et al. 2005*a*) the common bean genotype Quimbaya is equally Al-sensitive as VAX-1 after 4 h of Al treatment in spite of its higher CW negativity (Fig. 2c) and Al accumulation even after 8 h of Al treatment (Fig. 1b), indicating possible genotypic differences in Al compartmentation and/or binding in the root apex.

19 It has been argued that the strong binding of Al in the CW represents a detoxification 20 mechanism in squash (Le Van et al. 1994). However, the recovery from initial Al injury in 21 both genotypes during 4-8 h of Al treatment and after 24 h Al treatment in the Al-resistant 22 genotype Quimbaya were negatively correlated to the citrate non-exchangeable Al fraction of 23 the root apices (Fig. 4) and more specific to the stable-bound Al CW fraction (Fig. 8). This 24 suggests that the strong binding of Al to the pectic matrix of the CW is a main factor of Al 25 toxicity and not a resistance mechanism in common bean. In contrast to the stable-bound CW 26 Al fraction, there was no indication that the labile-bound (citrate-exchangeable) Al fraction

1 was related to Al-induced inhibition of root elongation (Fig. 8). This was unexpected, because 2 in maize this fraction appeared to contribute to explaining silicon (Si)-mediated amelioration 3 of Al toxicity (Wang et al. 2004). However, there seems to be a principal difference between 4 monocots and dicots in Al binding to CWs, which is not surprising given the difference in 5 CW composition (see above). This is well illustrated by the fact that treatment of CW with 50 6 mM BaCl₂ removed about 20% of the CW-bound Al in maize (Wang et al. 2004), and nearly 7 all Al adsorbed on wheat CW could be exchanged with 2.5 mM CaCl₂ (Zheng et al. 2004). In 8 contrast, BaCl₂ was unable to exchange any Al in common bean even after only short-term Al 9 treatment (Stass et al. 2007, this study). The significant negative relationship between root 10 elongation and citrate-exchangeable Al from intact root tips of genotype Quimbaya (Fig. 4a) 11 might be explained by the contribution of free apoplastic Al to this fraction (Fig. 8).

12 The fractionated extraction procedure allowed to separate operationally defined apoplastic 13 and symplastic Al fractions (Fig. 5). Among the 5 fractions the WFSF Al and the stable-14 bound CW Al-fractions are expected to best represent in vivo compartmentation of Al, the 15 first because it is recovered by centrifugation from the root tips without destroying the 16 compartmentation, the latter because it is expected to most slowly react during the extraction 17 steps. These two fractions showed a close negative relationship with root elongation-rate 18 reflecting recovery from initial Al stress particularly in genotype Quimbaya (Fig. 8). It is 19 difficult to decide whether the symplastic and the labile-bound CW Al fractions under or 20 overestimate the *in vivo* compartmentation. During the extraction process particularly during 21 the recovery of the cell sap, organic ligands may mobilize labile-bound CW Al or symplastic 22 Al is bound by CW due to a higher Al-binding strength of CW compared to symplastic 23 ligands (Rengel 1996). In spite of these uncertainties the fractionated extraction procedure has 24 proven to contribute to the understanding of Si amelioration of Al toxicity (Wang et al. 2004), 25 Si-accumulating and Si-excluding plant species in relation to their resistance against plant 26 pathogens (Heine et al. 2005, 2007), and Al accumulation of plant species like hydrangea

(*Hydrangea macrophylla* L.) and buckwheat (*Fagopyrum esculentum* Moench), which
 accumulate up to 70% Al in the symplast (B. Klug, personal communication) compared to 6 15% in common bean (Fig. 6).

4 The symplastic Al fraction neither reflect the recovery from initial Al stress in genotype 5 Quimbaya nor the enhanced Al sensitivity of VAX-1 after the temporary recovery period at 8 6 h Al treatment (Fig. 6). However, the trend of increasing symplastic Al contents with the 7 recovery and the significantly higher symplastic Al contents in Quimbaya compared to VAX-8 1 (Fig. 5) seems to indicate that transport of Al into the symplast is not a prerequisite for Al 9 toxicity. Higher symplastic Al contents may rather be indicative of enhanced/acquired Al 10 resistance which is in line with the observations by Vázquez et al. (1999) who ascribed 11 internalization of Al into the symplast contributing to Al tolerance in an Al-tolerant maize 12 genotype, and may indicate that also in common bean. Al internalization into 13 endosomal/vacuolar compartments may contribute to the recovery from initial Al stress as 14 reported for Arabidopsis (Illes et al. 2006). However, it is rather unlikely that this can explain 15 enhanced Al resistance because of the quantitatively small Al fraction in the symplast (Fig. 6). 16 The transitory (VAX-1) or sustained (Quimbaya) recovery from initial Al-induced inhibition 17 of root elongation (Fig. 1a) typical for pattern II plant species (see introduction) is related to a 18 decrease in Al contents of the root tip (Fig. 1b), particularly in the apical 2 mm region 19 (Rangel et al. 2007). The close negative correlation of root-elongation rate and Al contents of 20 the WFSF and the stable-bound CW fraction (Fig. 8) suggests that the recovery from initial Al 21 stress is related to the expression of an Al exclusion mechanism. This is in agreement with 22 previous studies indicating that citrate exudation is a mechanism of Al resistance in common 23 bean (Miyasaka et al. 1991, Mugai et al. 2000, Shen et al. 2002, Rangel and Horst 2006). 24 Evidence for the effects of organic acid secretion in Al resistance is substantial, but the mode 25 of action remains not well understood (Kinraide et al. 2005). Wehr et al. (2002) showed that 26 citrate and malate were able to remove Al from artificial Al-pectate gels suggesting that

1 exudation of organic acids would remove Al bound to pectin and this could alleviate toxicity. 2 However, the decrease of the Al content of the stable-bound CW Al fraction with increasing Al treatment duration as shown in Fig. 8 by root-released citrate appears to be improbable 3 4 because this fraction is defined as citrate non-exchangeable. It thus appears that once Al is 5 firmly bound it is unlikely to be released by the citrate exuded from the cells, unless the 6 citrate concentration in the apoplast is much higher than the concentration used for the 7 exchange (33 mM). Therefore, it is more likely that citrate released into the apoplast reduces 8 the binding of Al in the apoplast by complexing Al and decreasing the strength of Al binding, 9 thus preventing the strong binding of Al to the CW (Zheng et al. 2004). This allows resuming 10 cell division and cell elongation, and explains reduction of the Al contents in the root apex 11 through dilution by growth.

In conclusion, the results support the view that in common bean inhibition of root elongation cannot be explained by enhanced Al accumulation in the symplast. The present study indicates that the inhibition of root elongation is induced by apoplastic Al and that the induced and sustained recovery from the initial Al stress in the common bean genotype Quimbaya is mediated by reducing the stable-bound Al in the apoplast thus allowing cell elongation and division to resume

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1 Figure legends

2

3 Figure 1. Effect of Al treatment on the root-elongation rate (a) and total Al contents (b) of the 4 root tips of the common bean genotypes Quimbaya (Al-resistant) and VAX-1 (Al-sensitive) 5 grown in a simplified nutrient solution containing 0.5 mM CaCl₂, 0.5 mM KCl and 8 µM 6 H₃BO₃ without (Control) or with 20 μ M Al for up to 24 h, pH 4.5. Bars represent means ± SD, n = 4. For the ANOVA ***, ** denotes levels of significance at P < 0.001 and 0.01. 7 8 Means with the same letter are not significantly different between times within each genotype, 9 capital letters for Quimbaya and small letters for VAX-1; * denotes significant differences 10 between genotypes within each treatment time (Tukey test P < 0.05).

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12 Figure 2. Total cell-wall pectin-content (a) its degree of methylation (b) and unmethylated 13 pectin content (c) in 5 mm root tips of the common bean genotypes Quimbaya (Al-resistant) 14 and VAX-1 (Al-sensitive) grown in a simplified nutrient solution containing 0.5 mM CaCl₂, 15 0.5 mM KCl, and 8 µM H₃BO₃ without (Control) or with 20 µM Al for up to 24 h, pH 4.5. Bars represent means \pm SD, n = 4. For the ANOVA *, **, *** denote levels of significance at 16 17 P < 0.05, 0.01 and 0.001, n.s. = not significant. Means with the same letter are not 18 significantly different between times within each genotype, capital letters for Quimbaya and 19 small letters for VAX-1; * denotes significant differences between genotypes within each 20 treatment time (Tukey test P < 0.05).

1 Figure 3. Citrate-exchangeable (a) and non-exchangeable (b) Al contents in 5 mm root tips of 2 the common bean genotypes Quimbaya (Al-resistant) and VAX-1 (Al-sensitive) grown in a simplified nutrient solution containing 0.5 mM CaCl₂, 0.5 mM KCl, 8 µM H₃BO₃ and 20 µM 3 4 Al for up to 24 h, pH 4.5 Excised root tips were incubated for 15 min each, first in 50 mM 5 BaCl₂ and then in 33 mM Na₃Citrate. Bars are means \pm SD, n = 4. For the ANOVA *** 6 denotes a level of significance at P < 0.001. Means with the same letter are not significantly 7 different between times within each genotype, capital letters for Quimbaya and small letters 8 for VAX-1; * denotes significant differences between genotypes within each treatment time 9 (Tukey test P < 0.05).

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Figure 4. Relationship between root-elongation rate and citrate-exchangeable (a) or nonexchangeable (b) Al contents of root tips of the common bean genotypes Quimbaya (Alresistant) and VAX-1 (Al-sensitive) grown in a simplified nutrient solution containing 0.5 mM CaCl₂, 0.5 mM KCl, 8 μ M H₃BO₃ and 20 μ M Al for up to 24 h, pH 4.5. Excised root tips were incubated for 15 min each, first in 50 mM BaCl₂ and then in 33 mM Na₃Citrate. For the ANOVA *, *** denote levels of significance at P < 0.05 and 0.001.

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18 Figure 5. Aluminium contents of different cell compartments in 5 mm root tips of the 19 common bean genotypes Quimbaya (Al-resistant) and VAX-1 (Al-sensitive) grown in a 20 simplified nutrient solution containing 0.5 mM CaCl₂, 0.5 mM KCl, 8 µM H₃BO₃ and 20 µM Al for up to 24 h, pH 4.5. Bars represent means \pm SD, n = 4. For the ANOVA **, *** denote 21 22 levels of significance at P < 0.01 and 0.001. n.s. = not significant. Means with the same letter 23 are not significantly different between times within each genotype, capital letters for 24 Quimbaya and small letters for VAX-1; * denotes significant differences between genotypes 25 within each treatment time (Tukey test P < 0.05).

Figure 6. Relative distribution of Al contents of different cell compartments in the 5 mm root tips of the common bean genotypes Quimbaya (Al-resistant) and VAX-1 (Al-sensitive) grown in a simplified nutrient solution containing 0.5 mM CaCl₂, 0.5 mM KCl, 8 μ M H₃BO₃ and 20 μ M Al for up to 24 h, pH 4.5. The size of each pie chart represents the sums of all Al fractions. Means with the same letter are not significantly different between times in each fraction (Tukey test P < 0.05).

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Figure 7. Relationship between root-elongation rate and apoplastic (a) or symplastic (b) Al contents of root tips of the common bean genotypes Quimbaya (Al-resistant) and VAX-1 (Alsensitive) grown in a simplified nutrient solution containing 0.5 mM CaCl₂, 0.5 mM KCl, 8 μ M H₃BO₃ and 20 μ M Al for up to 24 h, pH 4.5. For the ANOVA *, *** denote levels of significance at P < 0.05 and 0.001.

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Figure 8. Relationship between root-elongation rate and the Al contents of three different apoplastic fractions in 5 mm root tips of genotype Quimbaya (Al-resistant) grown in a simplified nutrient solution containing 0.5 mM CaCl₂, 0.5 mM KCl, 8 μ M H₃BO₃ and 20 μ M Al for up to 24 h, pH 4.5. For the ANOVA *** denote levels of significance at P < 0.001.







3 Fig. 2

















3 Fig. 8