

# Four glycoside hydrolases are differentially modulated by auxins, cytokinins, abscisic acid and gibberellic acid in apple fruit callus cultures

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**Abstract**  $\alpha$ -L-Arabinofuranosidase,  $\alpha$ - and  $\beta$ -D-xylosidase, and  $\beta$ -D-glucosidase activity was detected in the soluble fraction (S-F) extracted with water and in the NaCl-released fraction (NaCl-F) extracted with a high-salt concentration buffer from apple callus cultures. The activity was found to be differentially modulated by the addition of various plant growth regulators (PGRs) to calluses that had lost their requirement for specific PGRs (“habituation” phenomenon).  $\alpha$ -L-Arabinofuranosidase activity was 93%, 130%, 126% and 186% higher in the NaCl-F from IAA-, IBA-, ABA- and GA<sub>3</sub>-treated callus than in that extracted from untreated callus while S-F  $\alpha$ -L-arabinofuranosidase activity was only 71%, 24%, 55% and 66% higher, respectively.  $\alpha$ -D-Xylosidase displayed low activity levels in both S-F and NaCl-F but 2iP-treated callus showed higher  $\alpha$ -D-xylosidase activity in both fractions than the control. 2,4-D

increased  $\alpha$ -D-xylosidase activity by 110% in the NaCl-F but decreased it by 40% in the S-F.  $\beta$ -D-Xylosidase activity increased by 99% in S-F from 2iP-treated callus but slightly decreased in the NaCl-F. In GA<sub>3</sub>-treated callus, NaCl-F  $\beta$ -D-xylosidase activity increased by 188%. S-F and NaCl-F from Picloram-treated callus showed undetectable or only slightly noticeable  $\alpha$ -L-arabinofuranosidase,  $\alpha$ -D-xylosidase and  $\beta$ -D-xylosidase activity. Interestingly,  $\beta$ -D-glucosidase activity rose 28-fold in the S-F extracted from Picloram-treated callus.  $\beta$ -D-glucosidase was the only enzyme assayed that greatly increased its NaCl-F activity after 10 subcultures, and the addition of any PGR to the callus culture –except for Picloram and ABA– decreased its activity, suggesting that this enzyme may be associated with certain stress conditions, such as PGR starvation or Picloram addition. This is the first report on glycoside hydrolases from fruit callus as modulated by different PGRs.

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## Abbreviations

ABA	Abscisic acid
$\alpha$ -Araf	$\alpha$ -L-Arabinofuranosidase
BAP	6-Benzylaminopurine
2,4-D	2,4-Dichlorophenoxyacetic acid
GA <sub>3</sub>	Gibberellic acid
$\beta$ -Glc	$\beta$ -D-Glucosidase

IAA	Indole-3-acetic acid
IBA	Indole-3-butyric acid
2iP	N <sup>6</sup> -(2-Isopentenyl)adenine
KIN	Kinetin
MS	Murashige and Skoog (1962) medium
NAA	1-Naphthaleneacetic acid
NaCl-F	Sodium chloride-released fraction
PGR	Plant growth regulator
PIC	4-Amino-3,5,6-trichloro-pyridine-2-carboxylic acid (Picloram)
PVPP	Polyvinylpyrrolidone
S-F	Soluble fraction
TDZ	Thidiazuron
$\alpha$ -Xyl	$\alpha$ -D-Xylosidase
$\beta$ -Xyl	$\beta$ -D-Xylosidase
ZEA	Zeatin

## Introduction

Glycoside hydrolases (EC 3.2.1.) are a very large and widespread group of enzymes (Davies et al. 2005) which hydrolyse the glycosidic bond between two or more carbohydrates or between a carbohydrate and a non-carbohydrate moiety. Although considerable advances have been made in unravelling the multiple functions of plant glycoside hydrolases, many other aspects are still unknown. Several glycoside hydrolases are bound to the cell wall where they may potentially trim terminal non-reducing glycosyl residues off various pectic and hemicellulosic homo- and heteropolysaccharides as well as different glycoconjugates (Rose et al. 2003; Sozzi 2004). Although these protein families may be vital for cell-wall remodelling, energy metabolism, signalling and defence during growth, development and stress responses (Farrokhi et al. 2006), the structural complexity of cell-wall polysaccharides makes it difficult to determine the specific sites at which these enzymes may cause functionally significant cell-wall changes. In addition, some glycoside hydrolases seem to be multifunctional enzymes. Conversely, other glycoside hydrolases are not bound to the cell wall, and may recognize moieties located in different molecules though their exact substrates and functions remain undefined.

Much of the existing research on glycoside hydrolase regulation has been performed on microorganisms, as they have potential industrial value in

the bioconversion of hemicelluloses and pectins. Yet, limited progress has been made in identifying plant growth regulators (PGRs) that modulate glycoside hydrolase activity in plant tissues since different problems have hindered research in this area. Studies with intact plant organs require vacuum infiltrations, sprays or dips of the organ in different PGR solutions. Therefore, entry and distribution of the PGRs are uncertain due to surface diffusion barriers and the treatment of each organ differs depending on its anatomy and morphology. For example, dipping fruits in concentrated PGR solutions can induce stress responses (Sozzi et al. 2000; Vendrell 1988). Another possible approach to test plant tissue responsiveness to physiological levels of different PGRs is the use of excised organs. The slicing of wild-type climacteric fruits causes wound-induced ethylene by excised discs but Sozzi et al. (2002) demonstrated the utility of a pericarp disc system from antisense ACC synthase fruit that excludes the interactions between applied PGRs and excision-stress ethylene biosynthesis.

Calyx and fruit cultures (Cohen 1996; Ishida et al. 1993) as well as callus cultures (Liu et al. 2006) have also been used to characterize different aspects of fruit metabolism. While a callus explant represents an artificial biological system, it provides a convenient easy-to-handle source of plant material that can be rapidly regenerated, and from which cell-wall enzymes may be obtained without difficulty. Also, callus cultures may allow comprehensive studies involving the addition of physiological levels of PGRs or other metabolites to the culture medium. However, to the best of our knowledge, there are no previous reports on the presence of glycoside hydrolases in apple fruit callus or on the modulation of their activity by different PGRs. The objective of this work was to characterize the activity of four glycoside hydrolases:  $\alpha$ -L-arabinofuranosidase ( $\alpha$ -Araf; EC 3.2.1.55),  $\alpha$ -D-xylosidase ( $\alpha$ -Xyl; EC 3.2.1.-),  $\beta$ -D-xylosidase ( $\beta$ -Xyl; EC 3.2.1.37) and  $\beta$ -D-glucosidase ( $\beta$ -Glc; EC 3.2.1.21) from apple callus, as affected by the addition of different PGRs.

## Materials and methods

### Origin of explants

Each explant consisted of a 5-mm apple-tissue cube obtained from the enlarged fleshy receptacle of a

75-day immature apple (*Malus × domestica* Borkh. 'Anna') grown in an orchard in Corrientes, Argentina.

### Culture initiation

The explants were surface sterilized by soaking for 1 min in 70% ethanol followed by immersion for 30 min in a solution of commercial bleach (containing 1.8% NaOCl, final concentration) and two drops of Tween<sup>®</sup>, and finally rinsed four times with sterile distilled water. Primary callus explants were placed individually in 11-ml glass tubes containing 3 ml of basal Murashige and Skoog (1962) medium (MS medium) supplemented with 1 mg l<sup>-1</sup> of 2,4-dichlorophenoxyacetic acid (2,4-D) and 0.1 mg l<sup>-1</sup> of 6-benzylaminopurine (BAP) as PGRs, and 3% sucrose as the energy source. The pH of each medium was adjusted to 5.8 with either KOH or HCl prior to the addition of 0.65% agar (Sigma, A-1296). Tubes were covered with aluminium foil and autoclaved at 0.101 MPa for 20 min.

### Physical culture conditions

The tubes were sealed with Resinite AF50<sup>®</sup> (Casco S.A.C. Company, Buenos Aires) and incubated at 27 ± 2°C in darkness. The callus showed neither morphogenesis nor organogenesis in that culture system.

To reduce any hormonal effect to a minimum, calluses were subcultured once (MS-1) or ten times (MS-10) at 30-days intervals onto a fresh basal regeneration medium (Murashige and Skoog 1962) supplemented with 3% sucrose as the energy source but with no PGR. Callus cultures can continue to grow in the basal medium with no PGRs due to the so-called "habituation" phenomenon. The term "habituation" has been given to all inheritable changes in the nutrient requirements (PGRs in particular) arising in cultured cell systems. Hence, when a normal culture or isolate loses its original requirement for auxins or cytokinins, it becomes "auxin- or cytokinin-habituated" (Hartmann et al. 1997). In some cases, a culture becomes habituated to more than one compound (Butcher 1973). This was the case with apple fruit tissue cultures that lose their requirement for 2,4-D and BAP and continue to proliferate even after the culture has been transferred to a medium without any PGR.

After ten subcultures, each PGR was added to the basal medium at a concentration of 5 mg l<sup>-1</sup>. The PGRs tested were the following: indole-3-acetic acid (IAA); two synthetic auxins [1-naphthaleneacetic acid (NAA) and indole-3-butyric acid (IBA)]; two auxinic herbicides [2,4-D and 4-amino-3,5,6-trichloro-pyridine-2-carboxylic acid (Picloram; PIC)]; three natural cytokinins [kinetin (KIN), zeatin (ZEA) and N<sup>6</sup>-(2-isopentenyl)adenine (2iP)]; two synthetic cytokinins [BAP and thidiazuron (TDZ)]; abscisic acid (ABA) and gibberellic acid (GA<sub>3</sub>). Three independent replicates per PGR treatment were prepared. After 30 d, actively growing calluses were harvested, frozen in liquid nitrogen and stored at -70°C until used. All chemicals were from Sigma Chemical Co. (St. Louis, MO, USA).

$\alpha$ -L-Arabinofuranosidase,  $\alpha$ - and  $\beta$ -D-xylosidase, and  $\beta$ -D-glucosidase activity

Glycoside hydrolases were extracted from 0.5 g of callus samples. Callus tissue was ground with 1 ml of cold sterile distilled water containing 1.5% (w/v) polyvinylpyrrolidone (PVPP). The suspension obtained was homogenised, centrifuged at 10,000g for 5 min at 4°C and the supernatant was called the soluble fraction (S-F). The procedure was repeated three times and each supernatant was checked for enzyme activity. The precipitate was then resuspended in 1 ml of 0.1 M sodium acetate, pH 5.5, containing 1.4 M NaCl. After 30 min at 4°C, the suspension was centrifuged at 10,000g for 15 min at 4°C, and the supernatant was called NaCl-released fraction (NaCl-F).

$\alpha$ -Araf,  $\alpha$ -Xyl,  $\beta$ -Xyl and  $\beta$ -Glc activity was measured using the corresponding *p*-nitrophenyl derivatives as substrates, namely: *p*-nitrophenyl- $\alpha$ -L-arabinofuranoside, *p*-nitrophenyl- $\alpha$ -D-xylopyranoside, *p*-nitrophenyl- $\beta$ -D-xylopyranoside and *p*-nitrophenyl- $\beta$ -D-glucopyranoside, respectively. The reaction mixture consisted of 125  $\mu$ l of 100 mM citrate buffer, pH 4.5, 100  $\mu$ l of 0.1% (w/v) bovine serum albumin (BSA), 100  $\mu$ l of enzyme solution (or an appropriate dilution) and 100  $\mu$ l of 13 mM substrate solution. After 1 h at 37°C, the reaction was stopped by addition of 425  $\mu$ l of 0.13 M Na<sub>2</sub>CO<sub>3</sub>. Blanks (0-time) were prepared by adding Na<sub>2</sub>CO<sub>3</sub> prior to the addition of substrate. Absorbance was measured at 400 nm. Free *p*-nitrophenol was used as standard. One unit of enzyme activity was defined as the amount of enzyme

that catalyses the liberation of  $1 \mu\text{g } p\text{-nitrophenol min}^{-1} \text{ mg}^{-1} \text{ protein}$ . Enzyme activity calculations were based on enzyme dilutions that gave a linear increase in free  $p\text{-nitrophenol}$  over a 2-h incubation period. Three independent replicates per PGR treatment were measured.

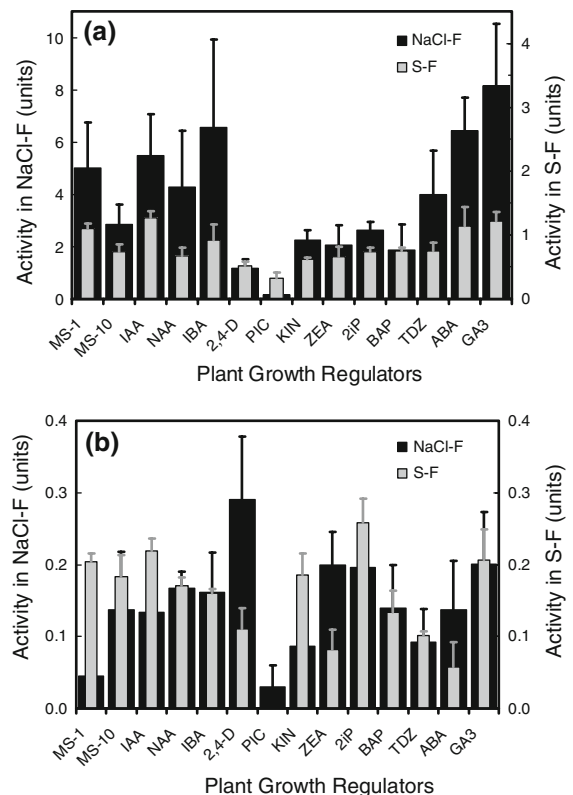
Protein concentration was assayed by the Coomassie Blue G dye-binding method using the Bio Rad reagent (Bio Rad, Richmond, CA, USA) and BSA as standard.

### Statistical analysis

Three independent replicates per PGR treatment were prepared and measured for each glycosidase activity and results were shown as the mean  $\pm$  SE of those three replicates. In addition, statistical significance was determined by one-way ANOVA using the PC-SAS software package (SAS Institute Inc., Cary, NC). The model assumptions of homogeneity of variance and normality were probed using Levene's test and Shapiro-Wilk's test, respectively. When these assumptions were not satisfied, data were transformed into ranks (Conover and Iman 1981) for further analysis. When a significant  $F$ -value was found, treatment means were compared using the Ryan-Einot-Gabriel-Welsch's test ( $P < 0.05$ ). This test controls the type I experimental error rate and has generally a lower type II error rate than the Tukey's studentized range test (Westfall et al. 1999).

### Results and discussion

$\alpha\text{-Araf}$ ,  $\beta\text{-Xyl}$  and  $\beta\text{-Glc}$  activities were mainly detected in the NaCl-F (Figs. 1a and 2). Most papers describe the separation of "cell-wall proteins" by tissue homogenisation in a low-ionic strength buffer or water (S-F) followed by successive washing of the cell-wall pellet with the same buffer to remove cytosolic proteins previous to extraction with a high-salt concentration buffer (NaCl-F) (Rose et al. 2003). Nevertheless, some positively-charged proteins could ionically bind to the cell walls once the plasma membrane has been disrupted because the polygalacturonate of cell-wall pectins can act as a polyanionic matrix. In this work, it is assumed that most of the activity obtained from homogenisation with strong saline buffer belongs to "cell-wall proteins" though



**Fig. 1**  $\alpha\text{-L-Arabinofuranosidase}$  (a) and  $\alpha\text{-D-xylosidase}$  (b) activity in the soluble (S-F) and the NaCl-released fraction (NaCl-F) from apple callus treated or not with different plant growth regulators. Each value represents the mean  $\pm$  SE of three replicates

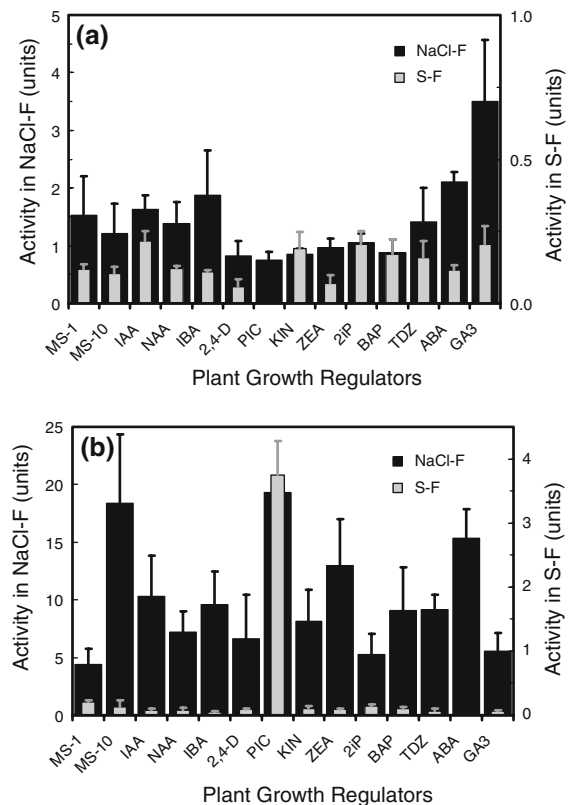
the final classification of these glycosidases depends on the verification of their subcellular localization by methods such as immunolocalization.

Callus subcultured ten times (MS-10) showed a decrease in NaCl-F  $\alpha\text{-Araf}$  activity (Fig. 1a). The addition of IAA, synthetic auxins, ABA or  $\text{GA}_3$  to the culture medium significantly increased NaCl-F  $\alpha\text{-Araf}$  activity, while PIC and, to a lesser extent, 2,4-D reduced the activity (Fig. 1a). NaCl-F  $\alpha\text{-Araf}$  activity from  $\text{GA}_3$ -treated callus was  $\sim 5135\%$  higher than that from PIC-treated callus. S-F  $\alpha\text{-Araf}$  activity from IAA-, ABA- or  $\text{GA}_3$ -treated callus was also greater than that from callus supplied with PIC (Fig. 1a). Itai et al. (2003) suggested that the *LeARF1* gene from tomato fruit is negatively regulated by ethylene while Sozzi et al. (2002) found three  $\alpha\text{-Araf}$  activities differentially modulated by synthetic auxins, gibberellic acid or ethylene, supporting the idea that total  $\alpha\text{-Araf}$  activity in tomato results from an

enzyme family whose members are responsive to different endogenous controls during growth and ripening.  $\alpha$ -Araf activity increases in ripening apples (Dick et al. 1990; Goulao et al. 2007; Yoshioka et al. 1995) while arabinose content decreases during the overripening stage (Peña and Carpita 2004). Nevertheless,  $\alpha$ -Araf is present not only during fruit ripening but also during the early stages of fruit development, both in apple (Goulao et al. 2007) and in Japanese pear, another pome fruit (Tateishi et al. 1996). Interestingly, the  $\alpha$ -Araf activity profile is consistent with the pattern of expression of *MdAF1* (Goulao et al. 2008) which displays high identity to *PpARF2* clone from Japanese pear (Tateishi et al. 2005). In callus,  $\alpha$ -Araf activity may be associated with the occurrence of arabinosyl residues in pectic side chains influencing cell-to-cell adhesion (Iwai et al. 2001).

In apple, almost no studies on  $\alpha$ -Xyl have been undertaken. In our experiments, S-F  $\alpha$ -Xyl activity was neither affected by successive callus subcultures (MS-10 vs. MS-1 activity; Fig. 1b) nor by addition of most of the PGRs (IAA, synthetic auxins, KIN, BAP, and GA<sub>3</sub>). S-F  $\alpha$ -Xyl activity decreased in 2,4-D-, ZEA-, TDZ- and ABA-treated callus, and was completely inhibited by PIC. On the other hand, NaCl-F  $\alpha$ -Xyl activity was greatly enhanced by callus subculturing (MS-10 vs. MS-1 activity; Fig. 1b), and only PIC restored low activity levels similar to those from MS-1 callus. NaCl-F  $\alpha$ -Xyl activity increased 5.4-fold upon treatment with 2,4-D, in comparison with that from MS-1 callus. Interestingly,  $\alpha$ -Xyl from 2,4-D-treated pea epicotyls was found to specifically cleave one of the  $\alpha$ -xylosidic linkages of xyloglucan-oligosaccharide substrates (O'Neill et al. 1989). Also,  $\alpha$ -Xyl role could be critical in that oligosaccharides lacking the xylosyl residue at the non-reducing terminus are probably unable to act as acceptors for xyloglucan endotransglycosylase (EC 2.4.1.207; Lorences and Fry 1993). Thus,  $\alpha$ -Xyl activity could diminish the presence of oligosaccharides vacant for transglycosylation during callus development.

The occurrence of  $\beta$ -Xyl activity has been verified in a range of fruit species (Sozzi 2004) but was not detected in apple using 4-methylumbelliferyl- $\beta$ -D-xyloside as substrate (Dick et al. 1990). In fact,  $\beta$ -Xyl activity against *p*-nitrophenyl- $\beta$ -D-xyloside was lower than  $\alpha$ -Araf and  $\beta$ -Glc in MS-1 callus (Figs. 1 and 2). Subculturing in the absence of PGRs did not affect S-F or NaCl-F  $\beta$ -Xyl activity levels (MS-10 vs. MS-1;



**Fig. 2**  $\beta$ -D-Xylosidase (a) and  $\beta$ -D-glucosidase (b) activity in the soluble (S-F) and the NaCl-released fraction (NaCl-F) from apple callus treated or not with different plant growth regulators. Each value represents the mean  $\pm$  SE of three replicates

Fig. 2a). Most of the PGRs assayed did not affect S-F  $\beta$ -Xyl activity. 2iP- and IAA-treated callus displayed higher S-F  $\beta$ -Xyl activity than 2,4-D- and ZEA-treated callus while activity was not noticeable upon treatment with PIC (Fig. 2a). In contrast, NaCl-F  $\beta$ -Xyl activity from GA<sub>3</sub>-treated callus increased  $\sim$ 188% while that from auxinic herbicide- or natural cytokinin-treated callus decreased 30 to 40% in comparison with that from MS-10 (Fig. 2a). The *PpAz152* transcript, which revealed high homology with plant  $\beta$ -Xyl, was found to be readily induced in peach by fruit ripening, leaf senescence and wounding but was also detected in early stages of fruit development and in flowers at all developmental stages (Ruperti et al. 2002). This transcript was not affected by the inhibition of ethylene perception. On the contrary, 1-methylcyclopropene partially counteracted the increase in  $\beta$ -Xyl activity during European pear ripening (Trincherro et al. 2004)



thus suggesting that this enzyme may be involved in a range of different physiological processes, modulated by species- and tissue-specific developmental and hormonal factors.

$\beta$ -Glc is also widely distributed in fruits (Sozzi 2004) including apple where only minor amounts of activity toward 4-methylumbelliferyl- $\beta$ -D-glucoside were found (Dick et al. 1990). S-F  $\beta$ -Glc activity was hardly noticeable in most cases and absent in ABA-treated callus (Fig. 2b). Interestingly, S-F  $\beta$ -Glc activity from PIC-treated callus rose 28-fold in comparison with that from MS-10 callus. In addition, NaCl-F  $\beta$ -Glc was the only enzyme assayed whose activity greatly increased after 10 subcultures (MS-10 vs. MS-1; Fig. 2b). Moreover, the addition of any PGR to the culture medium, except for PIC and ABA, decreased NaCl-F  $\beta$ -Glc activity in comparison with that from MS-10 callus. Thus, a  $\beta$ -Glc may be associated with certain stress conditions since the presence of PIC and/or PGR starvation (MS-10) positively modulates  $\beta$ -Glc activity. In fact, a gene homologous to a  $\beta$ -Glc was found to be expressed in apples exclusively during the infection process of the fruit by *Penicillium expansum* (Sánchez-Torres and González-Candelas 2003). To date,  $\beta$ -Glc has been detected in the cell suspension cultures of a range of species including carrot (Konno et al. 1996). In sweet cherry, the release of free glucose from cell walls by  $\beta$ -Glc suggests that this enzyme may play a role in cell-wall metabolism (Gerardi et al. 2001). In apple and other species,  $\beta$ -Glc may also be involved in the enzymatic hydrolysis of fruit glycoconjugates that play an important role as flavour precursors (Kramer et al. 1991; Schwab and Schreier 1988). Many flavour metabolites (e.g., shikimate-derived and terpenoid compounds) are typically conjugated to glucose as odourless  $\beta$ -D-glucopyranosides, or form more complex diglycosides with glucose being further conjugated with a second sugar unit (e.g.,  $\alpha$ -L-arabinofuranose,  $\beta$ -D-xylopyranose, etc.).

Because of the diverging activity profiles of the S-F and the NaCl-F in response to the addition of various PGRs to the callus cultures (Figs. 1 and 2), the existence of isoforms displaying different regulation ways should not be ruled out. On the other hand,  $\alpha$ -Araf,  $\beta$ -Xyl and  $\beta$ -Glc activity seem to be chiefly associated with the cell wall. A recent hypothesis states that three glycoside hydrolases ( $\beta$ -galactosidase,  $\beta$ -xylosidase and  $\beta$ -glucosidase) induced by

sugar starvation in *Arabidopsis* can remobilize sugars from cell walls (Lee et al. 2007). The activity of the glycoside hydrolases we studied seems to be differently modulated by auxins, cytokinins, abscisic acid and gibberellic acid. Thus, the possible addition of different PGRs may rule cell-wall metabolism in fruit callus cultures and will probably grant new opportunities to define cell-wall composition and quality in plant tissues. The exploitation of the so-called “habituation” phenomenon (fruit callus proliferating even after the culture has been transferred to a medium with no PGR) may be an interesting tool to understand the regulation of a plethora of genes and enzymes secreted into the cell wall.

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