

Phenylpropanoids, Phenylalanine Ammonia Lyase and Peroxidases in Elicitor-challenged Cassava (*Manihot esculenta*) Suspension Cells and Leaves

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Received: 12 September 2003 Returned for revision: 5 November 2003 Accepted: 13 February 2003 Published electronically: 14 May 2004

• **Background and aims** Control of diseases in the key tropical staple, cassava, is dependent on resistant genotypes, but the innate mechanisms are unknown. The aim was to study phenylpropanoids and associated enzymes as possible defence components.

• **Methods** Phenylalanine ammonia-lyase (PAL), phenylpropanoids and peroxidases (POD) were investigated in elicited cassava suspension cells and leaves. Yeast elicitor was the most effective of several microbial and endogenous elicitors. Fungitoxicity was determined against the cassava pathogens *Fusarium solani*, *F. oxysporum* and the saprotroph *Trichoderma harzianum*.

• **Key results** A single and rapid (≥ 2 –3 min) oxidative burst, measured as hydrogen peroxide, occurred in elicited cells. PAL activity was induced maximally at 15 h and was preceded by PAL mRNA accumulation, which peaked at 9 h. Symplasmic POD activity increased four-fold in cells, 48 h post-elicitation. POD isoforms (2–7 isoforms, pI 3.1–8.8) were detected in elicited and unelicited cells, extracellular medium and leaves but two extracellular isoforms were enhanced post-elicitation. Also expression of a cassava peroxidase gene *MecPOD1* increased in elicited cells. Only anionic forms oxidized scopoletin, with highest activity by isoform pI 3.6, present in all samples. Unidentified phenolics and possibly scopolin increased post-elicitation, but there was no enhancement of scopoletin, rutin or kaempferol-3-O-rutinoside concentration. Fungal germ tube elongation was inhibited more than germination by esculetin, ferulic acid, quercetin and scopoletin. *T. harzianum* was generally more sensitive than the pathogens and was inhibited by $\geq 50 \mu\text{g mL}^{-1}$ of ferulic acid and quercetin and $\geq 10 \mu\text{g mL}^{-1}$ of scopoletin.

• **Conclusions** Phenolic levels in cells were not enhanced and were, theoretically, too low to be inhibitory. However, in combination and when oxidized they may contribute to defence, because oxidation of esculetin and scopoletin by peroxidase and of esculetin by tyrosinase enhanced their fungitoxicity up to 20-fold.

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Key words: Cassava, *Manihot esculenta*, phenylalanine ammonia-lyase, peroxidase, oxidative burst, phenylpropanoids, kaempferol-3-rutinoside, rutin, scopoletin, scopolin, yeast elicitor, plant defence.

INTRODUCTION

Cassava (*Manihot esculenta*: Euphorbiaceae) is extensively cultivated in many tropical and sub-tropical countries and in some is the principal source of daily carbohydrates (Cock, 1985). Constraints on this staple crop include 25 different pathogens comprising a wide range of bacteria, viruses and fungi (Lozano, 1986). Usually, the only practicable and sustainable means of disease control is by deployment of resistant or tolerant genotypes. However, in contrast to many well-studied temperate crop or model species, the mechanisms of resistance in this plant to any pathogen have not been characterized. The interaction between elicitors of defence responses and plant cells has been much used as a tool to explore the potential resistance reactions to pathogens, but has not been exploited in cassava.

An almost ubiquitous feature of plant responses to incompatible pathogens or to elicitors is the activation of phenylpropanoid metabolism in which phenylalanine ammonia-lyase (PAL; EC 4.3.1.5) catalyses the first com-

mitted step of the core pathway of general phenylpropanoid metabolism. Branch pathways lead to the synthesis of compounds that have diverse functions in plants, notably in defence, such as cell wall strengthening and repair (e.g. lignin and suberin), antimicrobial activity (e.g. furanocoumarin, pterocarpan and isoflavonoid phytoalexins), and as signalling compounds such as salicylic acid (Hammerschmidt, 1999). The resulting phenolics are often converted into more reactive species by phenol oxidases and peroxidases (Mayer and Harel, 1979; Heath, 1980).

PAL genes can be regulated developmentally, induced by wounding, by low temperatures, by other stress conditions and by pathogen attack (Collinge and Slusarenko, 1987; Wu and Lin, 2002). PAL induction has been linked to defence responses that involve phenylpropanoids in numerous diseases. Typically, accumulation of PAL activity and mRNA is more rapid, higher and longer lasting in incompatible plant–pathogen interactions (Cui *et al.*, 1996). Inhibitors of PAL decrease resistance and associated changes in phenolic synthesis and incorporation into host cell walls, as in *Mla1* resistance of barley to powdery mildew and *Eucalyptus calophylla* to *Phytophthora cinnamomi* (Cahill and McComb, 1992; Zeyen *et al.*, 1995).

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Marked increases in PAL synthesis and corresponding mRNA occur in response to microbial or endogenous elicitors in many plant–pathogen systems (Sharan *et al.*, 1998). In particular, an elicitor derived from yeast induces PAL and the consequent accumulation of phytoalexins and other secondary metabolites in numerous plant species, including alfalfa, tobacco, *Lupinus albus*, apple, *Solanum khasianum* and soybean (Fahrendorf *et al.*, 1995; Mühlenbeck *et al.*, 1996; Wojtaszek *et al.*, 1997; Baier *et al.*, 1999; Borejsza-Wysocki *et al.*, 1999). This readily obtained, effective elicitor was deployed in this study.

Plant peroxidases (EC 1.11.17) are ubiquitous, heme-containing glycoproteins that catalyze oxidation of diverse organic and inorganic substances at the expense of hydrogen peroxide (H₂O₂). Higher plants have a number of peroxidase (POD) isoenzymes that are usually classified as anionic, neutral and cationic, based on their isoelectric points (Barz *et al.*, 1990). Typically there are 10–20 isoforms, some arising from divergent genes, which can differ by more than 50 % in peptide sequence, but some originate from post-translational modification of the same gene product (Zapata *et al.*, 1995; Lagrimini *et al.*, 1997). Peroxidase gene families are well described from many species, such as rice and parsley (Kawalleck *et al.*, 1995; Chittoor *et al.*, 1997). The increase of peroxidases after infection or elicitation often correlates with the appearance of new isoforms (Ludwig-Müller *et al.*, 1994; Adam *et al.*, 1995; Chittoor *et al.*, 1997). Anionic and neutral peroxidases are mainly cell wall-bound, and cationic forms are confined to the central vacuole (Perrey *et al.*, 1989; Kawalleck *et al.*, 1995), so the former are those mainly linked with defence because of their location. The roles in defence include oxidation of hydroxy-cinnamyl alcohols into free radical intermediates, phenol oxidation, cross-linking of polysaccharides and of extensin monomers, lignification and suberization (Chittoor *et al.*, 1997). Some forms can also generate active oxygen species as part of the oxidative burst during incompatible interactions (Adam *et al.*, 1995; Hiraga *et al.*, 2001). Increase in peroxidase activity during incompatible plant–pathogen/elicitor interactions is well documented and some peroxidases have been spatially and temporally associated with inhibition of pathogen growth (Adam *et al.*, 1995; Kawalleck *et al.*, 1995; Milosevic and Slusarenko, 1996; Chittoor *et al.*, 1997). Therefore, determination of peroxidase levels, isoforms and locations in cassava suspension cells and leaves challenged with yeast elicitor was undertaken to investigate their role as modifiers of the potential antimicrobial activity of cassava phenylpropanoids.

MATERIALS AND METHODS

Plant cell cultures and leaves

Stem stakes of cassava (*Manihot esculenta* Crantz) cultivar ‘MCol 22’ were provided by the Centro Internacional de Agricultura Tropical (CIAT) in Cali, Colombia, and were propagated in a glasshouse. The basal medium of Murashige and Skoog (MS) was used for the induction and maintenance of plantlets, callus and cell suspension cultures of

cassava. Cell suspension cultures were grown in (MS) medium, supplemented with the growth regulator 2,4-D (1 mg L⁻¹), kinetin (1 mg L⁻¹), plus 2 % glucose (Stamp and Henshaw, 1982). The cultures were incubated at 120 r.p.m. at 25 °C in darkness. Growth was determined by measurement of packed cell volume (pellet volume after centrifugation at 2500 g in 50 mL graduated conical centrifuge tubes).

Preparation of elicitors

Yeast cell wall glucan elicitor was prepared from baker’s yeast, autoclaved in citrate buffer, then the supernatant was precipitated with 50 % (v/v) (precipitate discarded), then 95 % (v/v) ethanol, and freeze-dried (Schumacher *et al.*, 1987). Cell wall glucan preparations from *Colletotrichum lindemuthianum* obtained by the method of Dixon and Lamb (1979) were kindly provided by R. Dixon (Samuel Roberts Noble Foundation, OK, USA) and G. P. Bolwell (Royal Holloway College, University of London, England). The carbohydrate content of glucan elicitor preparations was determined by the phenol–sulphuric acid method and concentrations are expressed as glucose equivalents (Dubois *et al.*, 1956). Oligogalacturonic acid elicitor was prepared by the method of Nothnagel *et al.* (1983) and generously donated by Dr F. Cervone (Dipartimento di Biologia Vegetale, Università di Roma ‘La Sapienza’, Italy) and Dr J. Dow (Purdue University, USA). Glutathione (GSH) was from Sigma (Poole, UK).

Treatment of cassava material with elicitors

Cassava suspension cells were treated with elicitors (water controls) after 5 d of subculture. After elicitation cells were harvested at time intervals, washed twice with 100 mL water on a porous-glass funnel with filter paper (Whatman No.1) then frozen in liquid nitrogen and stored at –80 °C. For some experiments, extracellular fluids were also analysed. Elicited cells to be analysed for production of peroxide were treated differently (see below). Leaf discs (1 cm diameter) were immersed in elicitor solution and incubated at 25 °C in the dark. Discs were collected at time intervals, frozen in liquid nitrogen and stored at –80 °C.

Enzyme extraction and PAL activity assay from cassava suspension cells

Crude enzyme extracts were prepared according to Hahlbrock and Ragg (1975), by homogenising 2 g of frozen cassava cells with 4 mL of 0.1 M sodium borate buffer pH 8.8 with an Ultra-turrax (IKA) homogenizer for 30 min at 0 °C; The slurry was centrifuged at 4 °C for 15 min at 20 000 g and the supernatant passed through pre-packed PD 10 columns (Sephadex G25 M). The reaction mixture consisted of 200 µL of the extract made up to 0.5 mL with 0.1 M sodium borate buffer (pH 8.8) and then added to an equal volume of the same buffer containing 0.02 M of L-phenylalanine or D-phenylalanine. The D-isomer is not degraded during the reaction and serves as a blank. The extinction coefficient of cinnamic acid is 16596 L mol⁻¹ cm⁻¹.

Incubation was at 30 °C, and A_{290} nm was measured after 0, 15, 30 and 60 min. One unit (U) of enzyme activity was defined as the amount of enzyme forming 1 pmol of *trans*-cinnamic acid from L-phenylalanine per min per mg of protein. The experiment was carried out in triplicate in three independent sets of experiments.

Enzyme extraction from cassava suspension cells and POD activity assay

Suspension cells (0.5 g) were ground for 15 min in a pestle and mortar on ice with 5 mL of 50 mM phosphate buffer, pH 6.0, 0.15 % polyvinylpyrrolidone (PVP, insoluble), 1 mM EDTA, 1 mM dithiothreitol (DTT), and 0.5 mM α -toluenesulfonyl fluoride 99 %. Debris was removed by centrifugation (4 °C, 45 min, 15 000 g). A standard curve was made with horseradish peroxidase (HRP) (Sigma, Type II). Peroxidase activity was assayed in the supernatant by the method of Van Gestelen *et al.* (1997). The 3.0 mL reaction volume contained 50 mM phosphate buffer (pH 6.0), 100 mM 3,5-dichloro-2-hydroxybenzenesulfonic acid, 10 mM 4-aminoantipyrine, and 150 μ L of enzyme extract (approx. 0.05 mg mL⁻¹ protein). After incubation at 25 °C for 5 min, the reaction was started by addition of 30 μ L of 100 mM H₂O₂, and the reaction followed at A_{510} nm. Three replicates were performed from three independent experiments.

Electrofocusing methods for POD isoforms

Symplasmic peroxidases. Extracts obtained as above from 3 g of cells were passed through a PD 10 column (Sephadex G25 M) then concentrated for about 2 h with polyethylene glycol 35 000 (PEG) in dialysis tubing to minimum volume and then dissolved in 100 μ L of 50 mM phosphate buffer pH 6.0 at 4 °C.

Apoplasmic peroxidases. After elicitation, cells were left for 15 min to sediment. 15 mL of extracellular fluids were collected into Falcon tubes by filtration through three layers of miracloth membrane, and then dialysed (Medicell International Size 2, internal diameter 18/32, molecular cut-off 12–14 kDa) for 2 d against excess 10 mM phosphate buffer, pH 6.0. The samples were concentrated for 6 h in dialysis tubing surrounded by PEG to minimum volume, then dissolved in 100 μ L of phosphate buffer at 4 °C.

Electrofocusing conditions for peroxidase isoforms. Peroxidase isoforms were separated over pH 3.5–9.5 in Ampholine PAG polyacrylamide gel by the method of Manchenko (1994). After running, the pI markers (pI 3–10) were cut from the gel and stained with Coomassie blue. The remainder of the gel was used for detection of POD isoforms by direct staining in 100 mL of 50 mM phosphate buffer (pH 6.0), with 100 mM 3,5-dichloro-2-hydroxybenzenesulfonic acid, 10 mM 4-aminoantipyrine and 100 mM H₂O₂ until red-brown bands appeared. Peroxidase isoforms with activity toward scopoletin were obtained by immersing

gels in 0.1 mM scopoletin dissolved in 0.1 M phosphate buffer, pH 6.0 (Gutiérrez *et al.*, 1995).

Protein determination

Protein was determined by the Bradford method (Stoscheck, 1990) with bovine serum albumin (BSA) as a standard.

Oxidative burst: H₂O₂ estimation

The method of Glazener *et al.* (1991) was used to estimate H₂O₂-scavenging by luminol (N-[4-aminobutyl]-N-ethylisoluminol)-dependent chemiluminescence in cassava suspension cells challenged with elicitor(s). Suspension cells were collected by centrifugation (200 g, 5 min) and washed with 150 mL of assay medium containing 0.5 mM of CaCl₂, 0.5 mM K₂SO₄, 175 mM mannitol and 0.5 mM MES (2-[N-morpholino] ethanesulfonic acid) buffer pH 5.8–6.0. Cell aggregates were removed by filtration through a 5 μ m pore plastic mesh then cells were suspended at 0.05 g mL⁻¹ of assay medium and preincubated for 2–3 h at 180 r.p.m. at 27 °C before use. Luminol-dependent chemiluminescence was measured with a luminometer (LUCY 1; Labtech) equipped with automatic sample dispensers. After addition of elicitors, 40 μ L of suspension cells were dispensed into black microtitre plates containing 100 μ L of 50 mM pH 9.0 CHES buffer (2-[N-cyclohexylamino] ethanesulfonic acid) and 60 μ L of peroxidase-luminol mixture was dispensed into each well to give final concentrations of 0.03 and 0.07 mg mL⁻¹, respectively. Total counts were measured over 30 s and converted to H₂O₂ equivalents.

Northern hybridization for PAL and POD transcripts

PAL (GenBank accession number AY036011) and POD (AY033386) cDNA probes were isolated from a post-harvest physiological deterioration library constructed in λ gt10 from roots of cassava cultivar 'Mnga 1' (Beeching *et al.*, 1997). PAL transcripts were identified by northern hybridization. The probe consisted of a 1.4 kb coding region of the PAL cDNA, excised by EcoRI, purified by agarose gel electrophoresis and labelled with α ³²P using an oligo-labelling kit (Pharmacia Biotech). The probe MecPOD1 consisted of a 600 bp coding region of the POD cDNA and was previously isolated in this laboratory. MecPOD1 shows highest similarity (up to 65 %) with adzuki bean (*Vincula angularis*) cationic peroxidase. Total RNA was extracted from cassava suspension cells, using the SV total RNA isolation system kit from Promega. Northern hybridization protocol was according to Sambrook *et al.* (1989). All lanes were loaded with 10 μ g of total RNA. The amount of RNA was determined spectrophotometrically based on the assumption that 1.0 A_{260} = 40 μ g mL⁻¹. All samples were adjusted to 1 μ g μ L⁻¹ RNA concentration. Equal loading of RNA was monitored. RNA markers were obtained from New England Biolabs and were stained with methylene blue according to Herrin and Schmidt (1988).

HPLC analysis of phenolics

Cassava cell suspensions were comminuted with a mortar and pestle in the presence of liquid nitrogen. 1 g of cells was suspended in 5 mL of ethanol 95 % (v/v), homogenized for 3 min in an Ultra-turrax (IKA), then incubated at 48 °C for 1 h. For cell suspension supernatants, 15 mL of medium was completely dried for 2 d in a speed-vacuum concentrator, and stored at -20 °C. 1 cm diameter discs from cassava leaves were ground with liquid nitrogen, then extracted with ethanol (1:10 w/v). Resulting suspensions were homogenized for 3 min then incubated at 48 °C for 1 h. Samples were centrifuged (5 min, 3000 g) then filtered (Whatman No.1), the supernatant recovered and evaporated to dryness at 40 °C. Samples, from cell suspensions, supernatants or leaf discs, were then prepared for HPLC analysis as detailed by Buschmann *et al.* (2000). Chlorophyll was removed from 200 µL samples in methanol by addition of 200 µL of water at 0 °C; chlorophyll precipitated after about 4 h, when samples were centrifuged (2 min, 4 °C, 10 000 g) then filtered (HPLC technology, PTFE 0.22 µm, 1.3 mm), transferred into brown glass sample vials and stored at -20 °C. Phenolics were identified by their retention times, by co-chromatography with reference compounds and by UV spectroscopy, as detailed by Buschmann *et al.* (2000). Quantification was based on reference compounds as external standards; standards were obtained from Sigma, except kaempferol 3-*O*-rutinoside (Roth Ltd, Germany).

Bioassays for antifungal activity

Fungal cultures. *Fusarium solani* and *F. oxysporum* from cassava were supplied by CIAT and cultured for 3 d at 25 °C on potato dextrose agar plates (PDA). *Trichoderma harzianum* was cultured for 4 d on malt agar plates. Conidial suspensions were made by flooding plates with sterile 1/6 strength Czapek Dox liquid medium pH 6 and adjusting to approx. 2.0×10^5 spores mL⁻¹.

Spore germination. Fungitoxicity of phenolic compounds was tested against spore germination on glass slides. Compounds were dissolved in absolute ethanol to give 10, 50, 100, 200, 500 and 1000 µg mL⁻¹. 50 µL of each solution was added to wells in Teflon™ microscope slides (BDH), evaporated, then 25 µL of spore suspensions were added. Conditions and assessment of germination and germ tube length were as described by Williams *et al.* (2002). In addition, crude extracts were tested by applying to thin layer chromatography (TLC) silica plates and determining toxicity to spores and to mycelial growth by spraying plates with fungal spores in nutrient solution; following incubation inhibitory zones were visualized as areas where mycelium failed to colonize the silica gel (Williams *et al.*, 2002). Phenolic solutions up to 1 mg mL⁻¹ were also added to sterile 13 mm antibiotic assay discs (Whatman, Maidstone, UK) and applied to newly established fungal cultures on agar plates; zones of inhibition would indicate toxicity to mycelial growth.

Fungitoxicity of oxidized phenolics. A preliminary survey was made of potential enzymic oxidation of major phenolics from cassava. Phenolics (250 µL of 100 µg mL⁻¹) in 60 mM phosphate buffer pH 6.0 were added to microtitre plates with hydrogen peroxide (at 0.1–1 % to establish optimum concentration) and 20 µL horseradish peroxidase (Sigma, Type II) or tyrosinase (Sigma), which was added without peroxide. Controls comprised combinations of all reagents except the phenolics. Plates were assessed visually and at A₄₂₀ nm to determine enzyme-mediated oxidation (evident as colour change).

Spores of *F. solani*, *F. oxysporum* and *T. harzianum* (100 µL of 2.0×10^5 spores mL⁻¹) were added to 50 µL mL⁻¹ phenolics (those shown previously to undergo oxidation) with 1:10 (v/v) peroxidase (plus H₂O₂ at 1, 0.05 or 0.02 % final concentrations) or tyrosinase, and after 30 min, centrifuged for 15 min at 2000 g. Spores were resuspended in 900 µL sterile solution of 15 % Czapek Dox, pH 6, and germination in 25 µL samples was determined on Teflon™ microscope slides as described above.

RESULTS

Evaluation of elicitor activity based on induction of the oxidative burst

A rapid, transient production of reactive oxygen species, is one of the earliest detectable responses to stress in plant cells (Bolwell, 1995), and is an indicator of the activity of potential elicitors. Several microbial and endogenous elicitors were tested for ability to induce production of extracellular H₂O₂. All elicitors were effective and caused a rapid and transient single burst starting after only 2–3 min with maximum production of 8 to >10 µM H₂O₂ after 5–35 min (Fig 1). Yeast cell wall glucan elicitor gave the fastest and highest response. It also induced other defence-related genes (Cooper *et al.*, 2001) and was used in subsequent experiments.

Elicitation of phenylalanine ammonia lyase activity and transcripts

A range of concentrations of yeast elicitor was used to elicit PAL mRNA accumulation in cell suspension cultures (Fig. 2). All concentrations were effective but 50 µg mL⁻¹ elicitor proved optimal and was used in subsequent experiments.

There was a progressive increase of PAL activity in cells after yeast elicitor treatment, with up to a four-fold increase relative to controls (Fig. 3). The maximum peak occurred at 15 h after elicitation. Northern blotting revealed PAL mRNA accumulated within 3 h and peaked at 9 h (Fig. 4). The overall profile over time mirrors that of changes in PAL activity, but the mRNA peak preceded that of enzyme activity by approx. 6 h, suggesting that the increases in PAL activity were due to expression of the corresponding PAL gene(s). No PAL expression was detected in elicited cassava leaf discs.

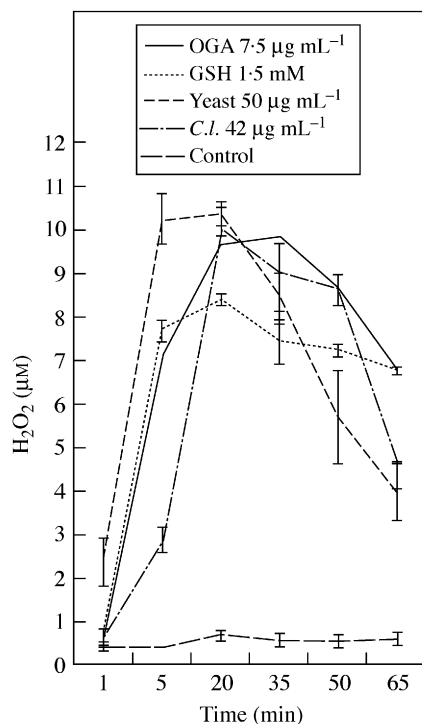


FIG. 1. Oxidative burst from cassava suspension cells challenged with microbial and endogenous elicitors. Extracellular hydrogen peroxide was measured by luminometry. OGA (oligogalacturonic acid $7.5 \mu\text{g mL}^{-1}$), GSH (glutathione 1.5 mM), *C.l.* (*Colletotrichum lindemuthianum* elicitor $42 \mu\text{g mL}^{-1}$), yeast (yeast elicitor $50 \mu\text{g mL}^{-1}$). Amounts of fungal elicitors are glucose equivalents. Error bars represent s.e. of three independent replicates.

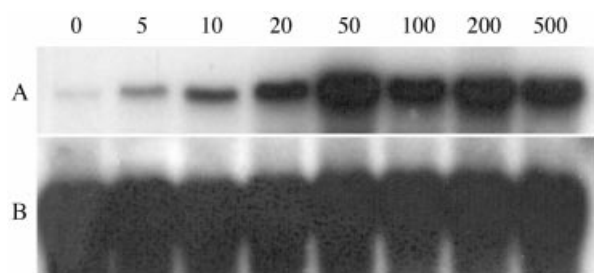


FIG. 2. Dose-dependent induction of PAL mRNA by yeast elicitor in cassava suspension cells. Northern blot of cassava cell suspension RNA purified 3 h after elicitation with $5\text{--}500 \mu\text{g mL}^{-1}$ glucose equivalents of glucan cell wall from yeast, and probed with the cassava PAL1 cDNA clone. (A) RNA from elicited cells probed with PAL cDNA; (B) the same RNA probed with 18S rDNA gene probe to check equal loading. RNA size-markers on the original gel indicated a size of approx. 2.5 kb for the PAL mRNA.

Identification and quantification of phenolic compounds in cassava cells and leaves

The induction of PAL in elicited cells suggested that phenolics might be produced during elicitation. This was

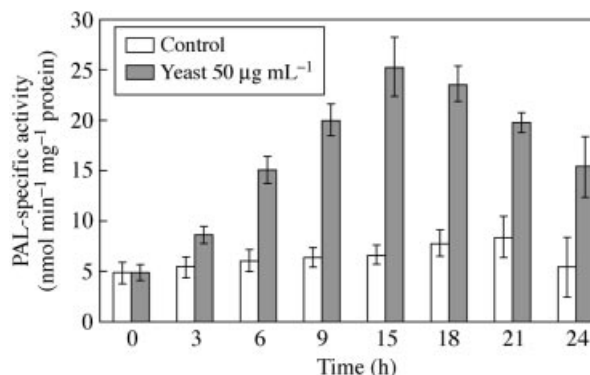


FIG. 3. Temporal induction of PAL in cassava suspension cells treated with yeast elicitor. Cassava cell suspensions (5 d growth) were treated with $50 \mu\text{g mL}^{-1}$ of glucan cell wall from yeast (solid bars) or control cells (open bars). Results are the means \pm s.e. of three replicates and are representative of three experiments.

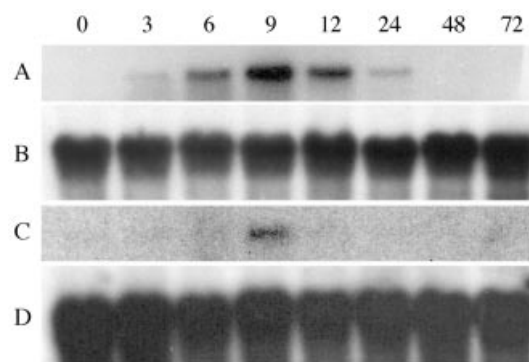


FIG. 4. Accumulation of PAL mRNA in elicited cassava cells. Northern blot of cassava cell suspension RNA purified over a time course after elicitation and probed with the cassava PAL1 cDNA clone. (A) RNA from cells elicited with $50 \mu\text{g mL}^{-1}$ of glucan cell wall from yeast, probed with PAL1 cDNA; (C) RNA from control cells probed with PAL1 cDNA. The same RNA probed with a 18S rDNA gene probe to serve as a loading control in: (B) elicited cells; and (D) in control cells.

investigated by HPLC of extracts from cassava cells, the medium in which they had been grown, and from leaves. Three phenolics, scopolin, rutin and kaempferol-3-*O*-rutinoside, were identified in the suspension cells. Scopolin was the only compound that sometimes increased on elicitation (Fig. 5A). In contrast, the medium in which the cells had been grown did not contain detectable phenolics (Fig. 5B). In cassava leaves there was a change in abundance of some compounds between elicited and control samples, but no change in the number of detectable compounds (Fig. 5C). Scopoletin in addition to scopolin, rutin and kaempferol-3-*O*-rutinoside were readily identifiable. However, a compound eluting at 6.5 min , which decreased on elicitation, and one eluting at 10.5 min , which increased, were not characterized.

Quantification of the four identified phenolics was made by HPLC with reference to pure samples according to Buschmann *et al.* (2000). Amounts were substantially

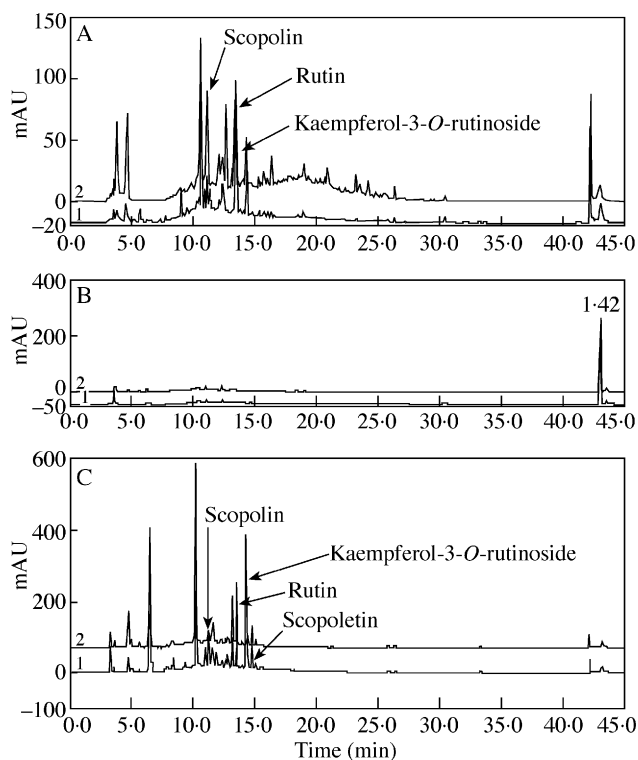


FIG. 5. HPLC detection of phenolics in elicited cassava cells and tissues. In all graphs the upper line corresponds to elicited cassava cells or leaves and the lower line to control cassava cells or leaves. (A) Elicited and sterile distilled water (SDW)-treated control cassava cells at 48 h; (B) elicited and SDW-treated medium at 48 h; and (C) elicited and SDW-treated cassava leaves at 48 h. Measurements at 280 nm.

greater in leaves than in cells and there was no consistent increase following elicitation. The maximum concentrations (nmol g^{-1} f. wt) found in elicited cells and leaves (bracketed figures) are: scopoletin, 0.07 (4.2); scopolin, 0.71 (6.0); rutin, 1.29 (9.6); and kaempferol-3-*O*-rutinoside 1.0 (20.5).

Antimicrobial activity of four major cassava phenolic compounds

Crude extracts from elicited and unelicited cassava cells showed fungitoxicity to spore germination by the slide bioassay and on the TLC plate bioassay with extracts equivalent to >50 mg f. wt mL^{-1} (data not shown). Therefore, the fungitoxicity of the phenolic compounds identified in the cassava leaves and cells (scopoletin and rutin) and phenolics previously identified in cassava roots undergoing post-harvest deterioration (esculetin, quercetin, ferulic acid; Buschmann *et al.*, 2000) was tested. Preliminary experiments ruled out the toxicity of rutin and kaempferol-3-*O*-rutinoside.

Figure 6 shows a typical dose-response curve for germ tube growth of *F. oxysporum*. Germ tube growth was reduced to 50% of the control value by 100–200 $\mu\text{g mL}^{-1}$ esculetin (0.6–1.2 μM), 50–100 $\mu\text{g mL}^{-1}$ ferulic acid (0.25–

0.5 μM), >1000 $\mu\text{g mL}^{-1}$ quercetin (3.3 μM) and 50–100 $\mu\text{g mL}^{-1}$ scopoletin (0.26–0.52 μM), but germination inhibition required >1 mg mL^{-1} of all phenolics (data not shown). Germ tube growth was generally more sensitive than germination and *T. harzianum* was usually more susceptible to phenolics than were the two pathogenic species. *Trichoderma harzianum* germ tube growth was inhibited by ≥ 50 $\mu\text{g mL}^{-1}$ ferulic acid and quercetin and ≥ 10 $\mu\text{g mL}^{-1}$ of scopoletin. None of the four phenolic compounds tested inhibited mycelial growth of the three fungi (data not shown).

Effect of oxidation on toxicity of phenolics

The major phenolics identified and tested above proved not to be highly antimicrobial. However, during infection and damage phenols and oxidases, such as peroxidase and tyrosinase, become decompartmentalized and could oxidize phenolic compounds to form antimicrobial quinones. In this study, peroxidase increased following elicitation (see below). Scopoletin and esculetin were readily oxidized by peroxidase and esculetin by tyrosinase *in vitro* (data not shown), as evident from the formation of coloured products. Therefore fungal spores were subjected to enzymatically oxidized esculetin and scopoletin (50 $\mu\text{g mL}^{-1}$).

With *T. harzianum*, inhibition was evident with scopoletin- and esculetin-oxidized derivatives; germination and germ tube growth were unaffected in all controls, including those containing hydrogen peroxide. In contrast to the native pure esculetin, which had no effect on germ tube growth at >1 mg mL^{-1} and required >100 $\mu\text{g mL}^{-1}$ to effect 50% inhibition of germination (Fig 6), following oxidation by peroxidase germination and germ tube growth were completely inhibited at 50 $\mu\text{g mL}^{-1}$; whereas, the product(s) resulting from tyrosinase activity reduced germination and germ tube growth to 28 and 19%, respectively. The quinone deriving from 50 $\mu\text{g mL}^{-1}$ scopoletin was more toxic to germination (reduced to 26%) than the native compound, which required >200 $\mu\text{g mL}^{-1}$ to produce half the effect (Fig 6). Similar trends occurred with the cassava pathogens but the degree of inhibition was less, with the oxidized forms again much more effective than the parent molecule, but typically resulting in 30–60% inhibition of germination and germ tube growth (data not shown).

Elicitation of peroxidase activity and transcripts

Yeast elicitor, triggered up to a four-fold increase of peroxidase activity in cells, which peaked at 48 h (Fig. 7). In contrast, peroxidase activity in the extracellular medium increased two-fold 12 h after elicitation and then decreased two-fold by 24 h (data not shown). Different peroxidase isoforms were revealed from these two locations and in leaves (Fig. 8). Suspension cells contained three or four POD isoforms, pls 3.1, 3.4, 3.5 and 8.8; those of pI 3.1 and 3.5 predominated. However, there was no discernible difference in either isoform number or their intensity between control and elicited cells. At least seven isoforms of pI (2.9, 3.0, 3.1, 3.4, 3.6, 8.8 and 8.8) were detected in the extracellular medium. Two isoforms (pI 8.8, but especially

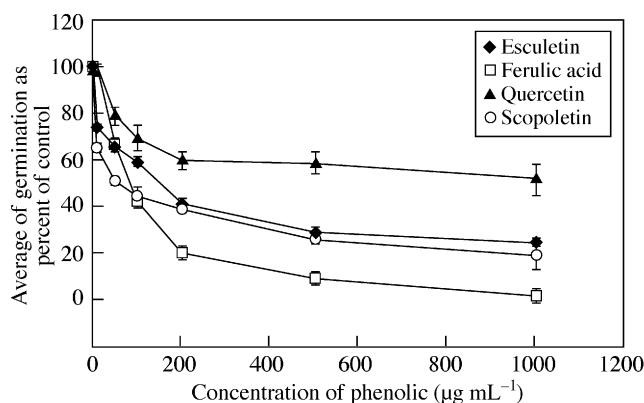


FIG. 6. Dose-response curve for germ tube growth of *F. oxysporum* treated with the four phenolic compounds.

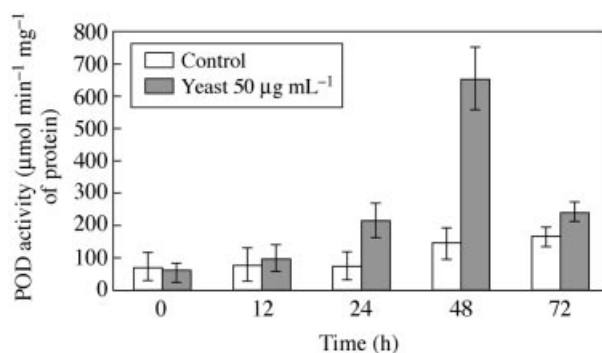


FIG. 7. Peroxidase activity in elicited and control cassava cells. Results are the means \pm s.e. of three replicates and representative of three experiments.

isoform pI 3.4) were enhanced in the elicited medium compared with the control. Two of the previously detected isoforms pIs, 3.4 and 3.6, were present in leaves and no difference was evident between the elicited and control leaves. Isoform pI 3.6 was present in all the samples.

Peroxidase isoforms from cassava suspension cells and leaves were tested for their capacity to oxidize scopoletin to a potentially more toxic form, as described earlier with a commercial peroxidase, by immersing isoelectric focusing gels with resolved POD isoforms in hydrogen peroxide and scopoletin (Fig. 9). All isoforms of horseradish peroxidase were able to oxidize scopoletin to a blue product (which later turned yellow), but only anionic peroxidases from cassava oxidized scopoletin. In particular, isoform pI 3.6, which was present in all samples including leaves, was highly effective in scopoletin oxidation.

Peroxidase changes during the time course of elicitation may derive from activation or from *de novo* synthesis of one of several isoforms. Therefore, it was of interest to investigate whether a representative corresponding gene (or genes) was activated. Total RNA was probed in northern blots with a cassava cDNA probe *MecPOD1*. Northern blots obtained with cassava cells after yeast elicitor challenge

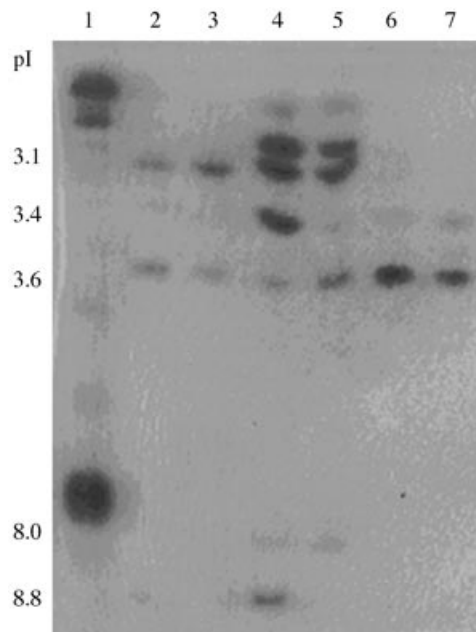


FIG. 8. Separation by isoelectric focussing of peroxidase isoforms of cassava cells, extracellular medium and leaves at 48 h after elicitation. Approximately 5 µg of cassava protein in 20 µL of extraction buffer was loaded into each lane. pI, isoelectric point. Lane 1, horseradish peroxidase II (positive control); Lane 2, cassava elicited cells; Lane 3, cassava control cells; Lane 4, extracellular medium from elicited cells; Lane 5, extracellular control medium; Lane 6, cassava elicited leaves; and Lane 7, cassava control leaves (SDW-treated).

showed that expression of *MecPOD1* started after 6 h and was maximal and similar after 12 and 24 h (Fig 10). No expression was observed in control cells, which shows that the non-induced activity is due to expression of peroxidase genes other than those coded by *MecPOD1*. Similarly, *MecPOD1* started to be induced 3 h after elicitation in leaves with maximal expression after 24 and 48 h.

DISCUSSION

A range of microbial and endogenous elicitors were tested for their ability to induce an oxidative burst in cassava suspension cells. All were effective but yeast elicitor always produced the most rapid and highest generation of H₂O₂, and of several other defence-related genes (data not shown). Therefore, yeast elicitor was chosen, rather than better-characterized but harder to obtain elicitors, to investigate phenylpropanoids and two associated enzymes.

Generation of H₂O₂ was rapid and in the form of a single burst, as was detected in soybean suspension cells treated with oligolacturonide elicitor (Chandra and Low, 1995). A second phase occurs following interactions with incompatible pathogens (Baker and Orlandi, 1995; Legendre *et al.*, 1993) and this was displayed by cassava cells when exposed to incompatible *Erwinia amylovora* (Cooper *et al.*, 2001). The peroxide produced may perform many defence-related functions (Baker and Orlandi, 1995), but in this context may

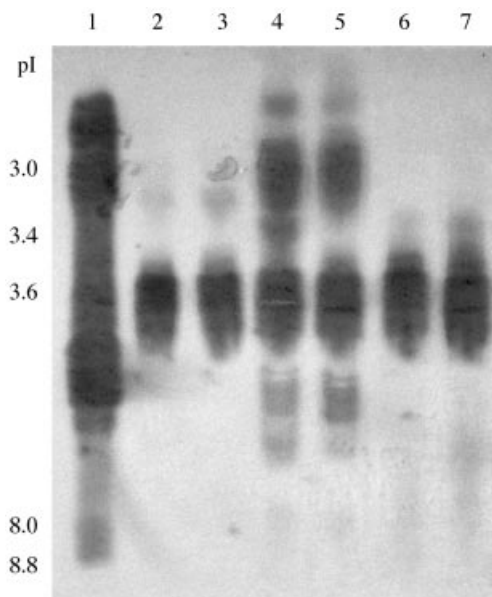


FIG. 9. Identification of cassava peroxidase isoforms with activity toward scopoletin. IEF gels were prepared as described for Fig. 8, then immersed in scopoletin as described in Materials and Methods. Coloured areas indicated formation of oxidized product corresponding to peroxidase isoforms.

contribute, as co-substrate, to the oxidation of phenolic compounds by peroxidase.

Maximal PAL activity in elicited cells occurred after 9 h. Similar kinetics have been found in cells and tissues of many resistant plant-pathogen interactions ranging from potato leaves and *Phytophthora infestans* to *Arabidopsis* suspension cells and elicitors (Fritzmeier *et al.*, 1987; Davis and Ausubel, 1989). Other members of gene families linked with phenylpropanoid biosynthesis may be activated later with different kinetics, including cinnamic acid 4-hydroxylase (C4H), chalcone isomerase (CHI), isoflavone reductase (IFR) and caffeic acid 3-*O*-methyltransferase (COMT) (Ni *et al.*, 1996).

No PAL expression was detected in elicited cassava leaves. Nevertheless, PAL activity and mRNA is induced in cassava leaves following infiltration with the incompatible plant pathogenic bacterium *Pseudomonas syringae* p.v. *phaseolicola* (R. Day and R. M. Cooper, unpubl. data), and Pereira *et al.* (1999) also obtained PAL activity and transcription of the PAL gene *MEPAL* in cassava leaves inoculated with incompatible *Xanthomonas cassavae*. It is feasible that there was insufficient exposure of leaf cells following immersion of leaf discs in elicitor solution.

Visible browning of elicited cassava cells may be indicative of enhanced phenylpropanoid metabolism but is more probably a consequence of phenolic oxidation. Browning of elicited suspension cells has often been noted (e.g. Fritzmeier *et al.*, 1987) but has sometimes been associated with cell death or decreased growth rate (Davis and Ausubel, 1989). However, elicitation did not decrease growth of challenged cassava suspension cells. Increased phenylpropanoid metabolism following infection or elicitor

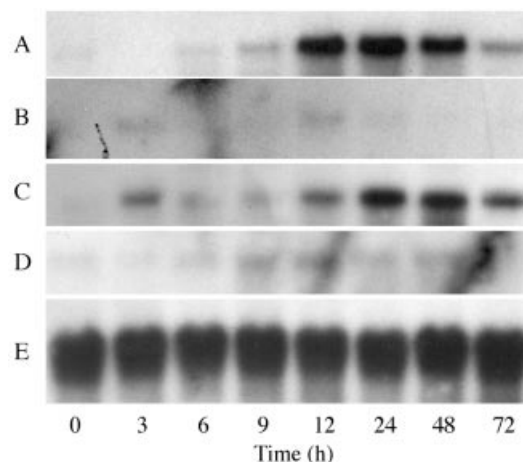


FIG. 10. Expression of the *MecPOD1* gene in cassava suspension cells and leaves after elicitation. Northern blots of cassava RNA purified after elicitation and probed with cassava *MecPOD1*. 10 µg of total RNA was loaded per lane. (A) POD in elicited cells; (B) POD in control cells; (C) POD in elicited leaves; (D) POD in control leaves; and (E) 18S rDNA in elicited leaves.

challenge has been found in many plants (Corchete *et al.*, 1993; Butland *et al.*, 1998). For example, alfalfa suspension cells challenged with yeast elicitor produced the isoflavonoid phytoalexin medicarpin (Dixon *et al.*, 1995). In other plants yeast elicitor has induced isoflavonoids and alkaloids, some of which were antimicrobial (Schumacher *et al.*, 1987; Borejsza-Wysocki *et al.*, 1999).

Phenolic compounds identified in cassava cells and leaves included the coumarin scopolin and its aglycone scopoletin, and the flavonoids kaempferol-3-*O*-rutinoside and rutin. Whilst some unidentified phenolics increased in elicited cells, none of these characterized phenolics, other than possibly scopolin, gave a clear increase after elicitation. This general pattern concurs with data of Davis and Ausubel (1989) and others (Jones, 1984) that PAL synthesis and enhanced phenolic accumulation are not invariably correlated. Phenolics were not detected by HPLC in the culture medium.

Some of these compounds and other phenolics have been reported in cassava to increase during wounding (scopoletin, scopolin, catechin, esculetin), post-harvest physiological deterioration (ferulic acid, esculetin, esculin, quercetin) or with defence responses to mealybug infestation (kaempferol-3-*O*-rutinoside, rutin) (Tanaka *et al.*, 1983; Wheatley and Schwabe, 1985; Calatayud *et al.*, 1994; Prawat *et al.*, 1995; Beeching *et al.*, 1999; Buschmann *et al.*, 2000). Scopoletin accumulates in many plants after infection and has been implicated as a phytoalexin in another member of the Euphorbiaceae, *Hevea brasiliensis* infected with *Microcyclus ulei* (García *et al.*, 1995) and with *Corynespora cassiicola* (Breton *et al.*, 1997). Confusingly, scopoletin has also been described as a phytoanticipin, in that it may be a defence compound mobilized from preformed precursors (Gutiérrez-Mellado *et al.*, 1996).

The toxicities of four available major phenolics from cassava were tested. The greater sensitivity of *T. harzianum* to the four phenolics than either *F. oxysporum* or *F. solani* may reflect that *T. harzianum* is a saprotroph, while the others are cassava-adapted pathogens. Tolerance to, or degradation of, antimicrobials from a host plant is often a key component of pathogenicity (Mansfield, 2000). Detoxification of scopoletin to scopolin, or by degradation, has been reported for several pathogens (García *et al.*, 1995).

Other studies have shown the fungitoxic effect of scopoletin (Ahl Goy *et al.*, 1993; García *et al.*, 1995; Breton *et al.*, 1997; Valle *et al.*, 1997). However, the efficacy of any antimicrobial agent depends upon it being at an appropriate concentration and at the right time and place *in vivo* (Williams *et al.*, 2002). Typically, compounds with a key role in resistance are active at $\leq 10^{-4}$ M (Mansfield, 2000), but many preformed phenolics do not possess this level of activity (Heath, 1980). Nevertheless, increase in scopoletin up to 35 nmol g⁻¹ f. wt and scopolin to 4 nmol g⁻¹ f. wt in leaves of the hybrid *Nicotiana glutinosa* × *Nicotiana debneyi* has been associated with resistance against *Cercospora nicotianae* and *Phytophthora parasitica* var. *nicotianae* (Goy *et al.*, 1993). Levels of the four identified phenolics in cassava cells were much lower than these reports; amounts were 0.07–1.29 nmol g⁻¹ f. wt in elicited cells and 4.2–20.5 nmol g⁻¹ f. wt in elicited leaves. These levels also contrast with the higher scopoletin levels of 34–124 nmol g⁻¹ f. wt in deteriorating cassava roots (Buschmann *et al.*, 2000). In addition, *H. brasiliensis* leaves accumulated up to 2 mM of scopoletin and this concentration was strongly inhibitory to germ tube elongation and conidial germination of leaf pathogens, including *Colletotrichum gloeosporioides* and *C. cassicola* (García *et al.*, 1995). Fliniaux *et al.* (1997) found that cell suspensions of different plant species vary greatly (0.001–5 mg g⁻¹) in the levels of scopolin and scopoletin they contain. Scopoletin has been reported to be inhibitory at approx. 2 mM to various fungi (Breton *et al.*, 1997; Valle *et al.*, 1997).

Critical information on fungitoxicity of the other phenolic compounds was not available; the few studies conducted with, for example, kaempferol, quercetin and rutin, used unstated levels or high and fixed concentrations, which does not allow for calculation of minimum inhibitory concentrations (e.g. El-Gammal and Mansour, 1986). However, as a generalization, phenolics in their unoxidized state are not highly antimicrobial (Heath, 1980). Therefore we investigated peroxidase, a defence-related enzyme with the capability of producing the more reactive, oxidized quinone forms.

Following elicitation, POD gene expression is often later than that of PAL, and for longer. Here, peak activity at 48 h followed maximum expression of a cassava peroxidase gene *MecPOD1* at 12–24 h in elicited cassava cells and leaves. In elicited *Arabidopsis* cells, POD mRNA was maximal at 24–48 h (Davis and Ausubel, 1989). Nevertheless, *MecPOD1* expression was first detectable much earlier after 3 h in leaves and 6 h in suspension cells. In elicitor-treated cassava suspension cells, symplasmic POD activity increased to a maximum at 48 h after elicitation.

Of the seven POD isoforms detected from cassava cells, anionic forms predominated and were mainly extracellular. Only two appeared to be up-regulated after elicitation, an isoform of pI 3.4 and, unusually, a cationic form of pI 8.8. POD genes from a family can be differentially regulated in response to various cues and in different tissues, as in rice and in bean (Adam *et al.*, 1995; Chittoor *et al.*, 1997). It is significant, perhaps, that only anionic isoforms in cassava oxidized one of the major cassava phenolics, scopoletin, and the most effective isoform pI 3.6 was present in suspension cells, leaf cells and extracellularly. Also, in tobacco suspension cells a particular isoperoxidase, A₃, was associated with the oxidation of scopoletin (Reigh *et al.*, 1975). However, the predicted amino acid sequence of the partial POD cDNA clone, MecPOD, used here in the northern analyses, has a theoretical pI and high homology with a cationic peroxidase. Therefore, it does not correspond to the anionic peroxidases with activity towards scopoletin. While POD isoforms were predominantly extracellular, as was the co-substrate H₂O₂, it is paradoxical that potential phenolic substrates for these enzymes were found mostly within the cells.

Furthermore, in cassava roots the increased peroxidase activity during PPD has been associated with scopoletin oxidation *in planta* (Wheatley and Schwabe, 1985). In these roots we found that mRNA for extensin (a cDNA clone MeHRGP1) showed a similar pattern of expression to that of MecPOD1 12–24 h after wounding (Han *et al.*, 2001). The cross-linking of extensin monomers by peroxidase may be yet another facet of the multi-component defences of cassava to pathogenic and biotic stress.

ACKNOWLEDGEMENTS

R.G.V. would like to thank the University of Bath, Department of Biology and Biochemistry, COLCIENCIAS (Colombian Science Council) and CIF (Centro Internacional de Física, Bogotá-Colombia), and R.D. acknowledges the ODA (now DFID) and the EC for support. This publication is in part an output from a research project (R8156 Crop Post-Harvest Programme) funded by the United Kingdom Department for International Development (DFID) for the benefit of developing countries. The views expressed are not necessarily those of DFID. We would like to thank Dr Kim Reilly for the cassava POD probe and Dr Han Yuanhuai for the PAL probe.

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