PISUM SATIVUM STRESS METABOLITES: TWO CINNAMYLPHENOLS AND A 2'-METHOXYCHALCONE

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Abstract—An HPLC study of copper(II) chloride treated *Pisum sativum* has shown that the cinnamylphenols obtustyrene [E-1-(4-hydroxy-2-methoxybenzyl)-2-phenylethylene] and xenognosin [E-1-(4-hydroxy-2-methoxybenzyl)-2-(4-hydroxyphenyl)ethylene] as well as the chalcone 4,4'-dihydroxy-2'-methoxychalcone accumulate as *de novo* metabolites in the stressed plant. This chalcone has not previously been identified from a plant source. The biosynthetic and stress response relationships of these compounds are discussed.

INTRODUCTION

A previous study [1] of the stress response of *Pisum* sativum L. has shown that most of the observed phenolic metabolites are related to the biosynthesis of the pterocarpinoid phytoalexins. However, three additional compounds are formed in response to copper(II) chloride stress which do not appear to fit the scheme of isoflavonoid biosynthesis. These compounds have been identified as the cinnamylphenols obtustyrene [E - 1 - (4 - hydroxy - 2 - methoxybenzyl) - 2 - phenylethylene], xenognosin <math>[E - 1 - (4 - hydroxy - 2 - methoxybenzyl) - 2 - methoxybenzyl) - 2 - (4 - hydroxyphenyl)ethylene] and 4,4' - dihydroxy - 2' - methoxychalcone. The biosynthetic and stress response relationships of these compounds are discussed.

RESULTS

A representative chromatogram of the phenolics which accumulate in stressed pea is given in Fig. 1. The peaks corresponding to the non-isoflavonoid phenolics, which are the subject of this study (1c, 2a, 2c) as well as the phytoalexin pisatin, have been identified on the chromatogram.

The characteristic UV spectrum [2] of metabolite 1c in combination with its molecular formula of $C_{16}H_{14}O_4$, which was derived from high resolution mass spectral measurements, suggested that the compound was a dihydroxymethoxychalcone. The

mass spectrum had a hydroxy substituted cinnamovl derived fragment and a hydroxy, methoxy substituted benzoyl derived fragment. The ¹H NMR aromatic couplings indicated that the benzoyl part of the molecule was 2,4-substituted while the cinnamoyl part was 4-substituted. Thus, the isolate was either 4,4' - dihydroxy - 2' - methoxychalcone (1c) or 2',4 dihydroxy - 4' - methoxychalcone (1d). The identification of the metabolite as the 2'-methoxy isomer (1c) was based on the expected [2] effect of aluminium chloride on the UV spectrum of a 2'hydroxychalcone and a comparison of its chromatographic properties with synthetic samples of both the 2'-hydroxy and the 2'-methoxychalcones. The synthesis of the 2'-hydroxy isomer followed a known aldol condensation route [3]. Unfortunately, it was



Fig. 1. A representative chromatogram of the Pisum sativum stress metabolites.

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Fig. 2. Flavonid-isoflavonoid and cinnamylphenol biosynthesis. Potential interconnection (----).



Fig. 3. Probable route to the formation of a quinonemethide from xenognosin.

found that the condensation of 4' - hydroxy - 2' - methoxyacetophenone with 4 - hydroxybenzaldehyde gave a poor yield of impure 4,4' - dihydroxy - 2' -

methoxychalcone. However, this compound was obtained in good yield and purity by the sequential methylation and debenzylation of 4,4'-bis(benzyloxy)-2'-hydroxychalcone.

Metabolite 2a had a high resolution molecular formula, determined by mass spectroscopy, of C₁₆H₁₆O₃. An ABX₂ coupling pattern in the ¹H NMR spectrum and an increase in mass of 28 after reaction with methyl iodide-potassium carbonate indicated that the compound did not contain a carbonyl oxygen and was therefore a dihydroxymethoxycinnamylphenol. Analysis of the mass spectrum, which had the tropylium fragments C₇H₇O and C₈H₉O₂, in combination with the similarity of the 'H NMR aromatic couplings of this isolate and the previously identified chalcone indicated that the phenyl substituent patterns were 4-hydroxy and either 4-hydroxy-2methoxy or 2-hydroxy-4-methoxy. The position of the double bond was determined by HPLC analysis of the products obtained from the ruthenium tetraoxide oxidation [4] of the isolate. This degradative procedure, which gave 4-hydroxybenzaldehyde as the only observed product, proved that the isolate was



either E - 1 - (4 - hydroxy - 2 - methoxybenzyl) - 2 - (4 - hydroxyphenyl)ethylene (2a) or its 2 - hydroxy - 4 - methoxybenzyl isomer (2b). Chromatographic comparison of the isolate with synthetic samples of these compounds proved that it was the 4 - hydroxy - 2 - methoxybenzyl isomer (2a). This isolate is a rare natural product recently characterized for the first time from a exudate of Astragalus species and named xerognosin [5].

Metabolite 2c was identified as obtustyrene [E - 1 - (4 - hydroxy - 2 - methoxybenzyl) - 2 - phenylethylene], which has been previously identified from*Dalbergia retusa*by Gregson*et al.*[6] by methods analogous to those used for 2a.

DISCUSSION

The general scheme of flavonoid/isoflavonoid biosynthesis is given in Fig. 2(A). As illustrated, the formation of the isoflavone pyrane ring requires a free 2'-hydroxyl on the chalcone [7]. Thus the 2'methoxychalcone does not appear to be an intermediate in flavonoid/isoflavonoid synthesis.

The cinnamylphenols have previously only been isolated from heartwoods. Consequently, the currently favoured route for cinnamylphenol biosynthesis, which is based on cinnamylphenol/3,3-diarylpropene co-occurrence in some plants and the *in vitro* observation of diarylpropene and cinnamylphenol formation in the condensation of cinnamyl alcohols with phenols [8,9], requires that a cinnamyl pyrophosphate condense with a phenol (or its equivalent) to give a 3,3-diarylpropene and/or a cinnamylphenol [8] (Fig. 2B). There is no evidence to indicate that the cinnamylphenols and the chalcones are biosynthetically related. However, the structural similarity of the 2'methoxychalcone and the cinnamylphenol xenognosin suggests that they may be interconnected. The identification of both of these compounds as *de novo* metabolites of *P. sativum* provides a convenient system for further studies on their biosynthesis.

The role of these compounds in the plant's stress response is not immediately apparent because if they are antifungal it is likely that they would have been observed by Cruickshank and Perrin [10] in their initial TLC bioassays of stressed pea metabolites. However, we are currently investigating the possibility that xenognosin (2a) may function in the plant's disease response through its anticipated ability to form reactive quinone-methides [10] (Fig. 3). These compounds could participate in the resistance process by reaction with proteins and enzymes or by polymerization to tannins and lignans which can act as protein precipitants or as physical barriers to parasite expansion [11].

EXPERIMENTAL

The HPLC system and UV analysis procedures used in this study have been described previously [12]. The procedures for growth and stress of the pea plants (*Pisum* sativum cv Melting Sugar) have also been described previously [1]. MS (direct insertion probe; ionization voltage 70 eV). Chromatographic solvents were: petrol (PE), Et₂O (E), CH₂Cl₂ (MDC) and Me₂CO (A). The metabolites were isolated by HPLC as described previously [12]. Obtustyrene was identified by direct comparison of its HPLC, TLC, UV and MS characteristics with a synthetic sample [6].

4,4' - Dihydroxy - 2' - methoxychalcone (1c). Identification. The HPLC isolate was further purified by TLC on Si gel [PE-E (1:2); $R_f = 0.38$]. Analysis of the resultant sample gave UV λ^{MeOH} nm: 253, 349; NaOMe 257, 419; NaOAc 255sh, 358sh, 393; NaOAc/H3BO3 351; AlCl3 235, 350; AlCl₃/HCl 235, 350; ¹H NMR [100 MHz, (CD₃)₂CO]: δ 7.50-7.64 (4H, m, C- β , C-2, C-6, C-6'); 7.28 (1H, d, J = 18 Hz, C- α); 6.91 (2H, d, J = 8 Hz, C-3, C-5); 6.59 (1H, d, J = 2 Hz, C-3'); 6.52 (1H, dd, J = 2,8 Hz C-5'); 3.92 (3H, s, -OMe); MS: Found: $[M]^+$ (HRMS), 270.0885. $C_{16}H_{14}O_4$ requires: $[M]^+$, 280.0892. m/z (rel. int.), 270 $[M]^+$ (100), 269 (23), 255 (38), 242 (31), 164 (38), 163 (18), 151 (80), 147 (33), 119 (25), 107 (89). Synthesis. 4,4'-Bis(benzyloxy) - 2' - hydroxychalcone [13] (1a) was methylated with MeI-K₂CO₃ to give 4,4' bis(benzyloxy) - 2' - methoxychalcone (1b). The product was crystallized from Me₂CO-EtOH (5 mM scale; 82% yield). mp 94-95°, Found: C, 79.43; H, 5.85; [M]⁺ (HRMS), 450.1808. $C_{29}H_{26}O_4$ requires: C, 79.42; H, 5.98; [M]⁺, 450.1831; UV $\lambda_{max}^{CH_2Cl_2}$ nm (log ϵ): 236 (3.26), 340 (3.48); IR $\nu_{\text{max}}^{\text{CH}_2\text{Cl}_2}$ cm⁻¹ 3001, 1601, 1250, 1163, 1129, 1020, 830; ¹H NMR (100 MHz, CDCl₃): δ 7.18-7.76 (15H, m, C-α, C-β, C-2, C-6, $2 \times s$, -CH ϕ); 3.78 (3H, s, OMe); MS m/z (rel. int.) 450 [M]⁺, (14), 359 (12), 197 (12), 92 (15), 91 (100).

The benzyloxychalcone (230 mg, 0.510 mmol) was debenzylated in conc. HCl-glacial HOAc (2:3, 25 ml) by stirring the reaction mixture for 2 hr in a H₂O bath at 48°. The product was extracted into Et₂O after the addition of H₂O (100 ml) to the reaction mixture. The Et₂O layer was successively extracted with H₂O, 5% aq NaHCO₃ and H₂O. After drying over dry Na₂SO₄ and filtration the solvent was removed on a rotary evaporator. The residue was chromatographed on Si gel (15 g, 20×1 cm column) with MDC-A (4:1) and the resultant solid crystallized from aq. EtOH to give 70 mg (51%) of 4,4' - dihydroxy - 2' - methoxychalcone (1c) as orange microcrystals. This product was found to be identical with the isolate by HPLC, TLC, UV, ¹H NMR and MS. In addition it gave mp 210-212°, Found: C, 70.90; H, 5.11. C₁₆H₁₄O₄ requires C, 71.10; H, 5.22; UV λ_{max}^{MeX} nm (log ϵ): 234 (4.00), 349 (4.25); IR ν_{max}^{nujol} cm⁻¹: 1560 br, 1330, 1315, 1250 br, 1218 br, 1170, 1120, 825.

E - 1 - (4 - Hydroxy - 2 - methoxybenzyl) - 2 - (4 - 4)hydroxyphenyl)ethylene (2a) (xenognosin). Identification. The HPLC isolate was further purified by Si gel TLC (PE-E (1:1); $R_f = 0.58$). Analysis of the resultant sample gave UV λ_{max}^{MeOH} nm: 261.5; NaOMe 281; NaOAc, NaOAc-H₃BO₃, AlCl₃ and AlCl₃-HCl 261.5; ¹H NMR [100 MHz, (CD₃)₂CO]: δ 7.22 (2H, d, J = 9 Hz, C-2, C-6); 6.97 (1H, d, J = 8 Hz, C-6'); 6.76 (2H, d, J = 9 Hz, C-3, C-5); 6.38-6.50 (2H, m, C-3', C-5'); 6.16–6.32 (2H, m, C- α , C- β); 3.80 (3H, s, OMe); 3.36 $(2H, d, J = 6 Hz, C-X \text{ of } ABX_2); MS. Found: [M]^+ (HRMS),$ 256.1105. $C_{16}H_{16}O_3$ requires: [M]⁺, 256.1100; UV λ_{max}^{MeOH} nm: 256 m/z (rel. int.) [M]⁺ (100), 255 (19), 241 (17), 239 (17), 225 (21), 161 (11), 150 (15), 149 (23), 137 (63), 133 (26), 132 (23), 131 (41), 122 (36), 121 (61), 107 (56). Synthesis. p-Hydroxycinnamyl alcohol (0.750 g, 5 mmol) was dissolved with m-methoxyphenol (1.24 g, 10 mmol) in 20 ml 75% HOAc-H₂O and stirred for 15 min at 60°. The products were extracted into Et₂O after the addition of 100 ml H₂O. The Et₂O layer was extracted successively with H₂O, 5% aq. NaHCO₃ and H₂O followed by drying, filtration and removal of the solvent on a rotary evaporator. Chromatography on Sigel $(100 \text{ g}, 45 \times 4 \text{ cm})$ using PE-E (1:1) gave 0.44 g of the mixed isomers 2a and 2b. Separation of these isomers by CC as above using PE-E (2:1) and MCD-EtOH (19:1) gave 190 mg of the faster moving and 205 mg of the slower moving isomers. Both isomers resisted all attempts at crystallization (though both were pure as measured by TLC) and both proved to be too unstable for satisfactory microanalysis. However, both isomers, when reacted with MeI in the usual manner, gave the expected trimethoxy product (2e). The second isomer to be eluted from the column was found by HPLC and TLC to correspond to the plant isolate. Its identification as 2a was confirmed by TLC and HPLC comparison with the product obtained from the reduction of chalcone 1c with AlH₃ [14]. The synthetic product was also found to be identical with the isolate by UV, ¹H NMR and MS. Found: [M]⁺ (HRMS), 256.1105. $C_{16}H_{16}O_3$ requires: [M]⁺, 256.1100; UV λ_{max}^{MeOH} nm: $(\log \epsilon)$ 261 (4.0); ¹H NMR [270 MHz, (CD₃)₂CO]: δ 7.22 (2H, d, J = 9 Hz, C-2, C-6); 6.96 (2H, d, J = 8 Hz, C-6'); 6.76 (2H, d, J = 9 Hz, C-3, C-5); 6.45 (1H, d, J = 2 Hz, C-3'); 6.38 (1H, *dd*, J = 2,8 Hz, C-5'); 6.33 (1H, *d*, J = 16 Hz, C- β); 6.16 (1H, $td, J = 6,16 \text{ Hz}, \text{ C-}\alpha$; 3.82 (3H, s, OMe); 3.36 (2H, d, J = 6 Hz, C-X of ABX₂).

Synthesis (2e). p-Hydroxycinnamyl alcohol (0.750 g, 5 mmol) was dissolved with resorcinol (1.10 g, 10 mmol) in 50 ml 75% HOAc-H₂O and stirred for 15 min at 70°. The product was isolated as above. CC purification of the residue on Si gel [100 g, 45 × 4 cm column, PE-E (1:3)] gave 0.449 g (37%) of 2d as a glass. Found: [M]⁺ (HRMS), 242.0925. C₁₅H₁₄O₃ requires: [M]⁺, 242.0943; UV λ_{max}^{meOH} (log ϵ): 262 (4.20); NaOMe 240 sh, 283; ¹H NMR [270 MHz (CD₃)₂CO]: δ 7.17 (2H, d, J = 9 Hz, C-2, C-6); 6.87 (2H, d, J = 8 Hz, C-6'); 6.72 (2H, d, J = 9 Hz, C-3, C-5); 6.37 (1H, d, dJ = 2 Hz, C-3'); 6.23-6.33 (2H, m, C- β , C-5'); 6.16 (1H, td, J = 6,16 Hz, C- α); 3.37 (2H, d, J = 6 Hz, C-X of ABX₂); MS 242 [M]⁺, (100), 241(10), 225(17), 149(10), 147(40), 135(10), 133(32), 132(13), 131(28), 123(42), 119(29), 107(69). The trihydroxy product (108 mg, 0.44 mmol) was methylated with MeI-K₂CO₃. The product was purified by TLC on Si gel [2000 μ m layer, 40 × 40 cm plate, PE-E (4:1), $R_f = 0.75$] and crystallized from aq. MeOH to give 72 mg (57%) of 2e as white plates, mp 66-67°. Found: C, 75.60; H, 6.82; [M]⁺, 284.1432. C₁₈H₂₀O₃ requires: C, 76.03; H, 7.09; [M]⁺, 284.1412. UV $\lambda_{\max}^{CH_2Cl_2}$ nm (log ϵ): 262 (4.35); IR $\nu_{\max}^{CHCl_3}$ cm⁻¹: 1611, 1506, 1465 br, 1296, 1252 br, 1184, 1165, 1127, 1045, 975, 845; ¹H NMR [270 MHz, (CD₃)₂CO]: δ 7.26 (2H, d, J = 9 Hz, C-3, C-5); 6.44 (1H, d, J = 2 Hz, C-3'); 6.41 (1H, dd, J = 2, 8 Hz, C-5'); 6.33 (1H, d, J = 16 Hz, C- β); 6.19 (1H, td, J = 6 Hz, C- α); 3.80; 3.76; 3.75 (9H, 3×s, OMe); 3.43 $(2H, d, J = 6 Hz; C-X \text{ of } ABX_2); MS 283 [M]^+ (100), 282(13),$ 269(11), 253(58), 151(18), 146(20), 145(42), 121(39).

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