

AN ABSTRACT OF THE THESIS OF

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The metabolism of [8-14C]zeatin was examined in embryos of Phaseolus vulgaris cv. Great Northern (GN) and P. lunatus cv. Kingston (K) in an attempt to detect genetic variations in organized plant tissues. Metabolites were fractionated by HPLC, and identified by chemical and enzymatic tests and GC-MS analyses. Five major metabolites were recovered from P. vulgaris embryo extracts, ribosylzeatin, ribosylzeatin 5'-monophosphate, the O-glucoside of ribosylzeatin and two novel metabolites, designated as I and II. Based on results of degradation tests and GC-MS analyses, I and II were tentatively identified as O-ribosides of zeatin and ribosylzeatin. In embryos of P. lunatus, however, metabolites I and II were not present. The major metabolites were ribosylzeatin, ribosylzeatin 5'-monophosphate and the O-glucosides of zeatin and ribosylzeatin. The zeatin metabolites recovered were the same for embryos of different sizes but their quantities varied with embryo size and incubation time. The genetic differences appear to be embryo-specific and may

be useful in the studies of the possible relationship between abnormal interspecific hybrid embryo growth and hormonal derangement in Phaseolus. In addition, analyses of both organized (intact) and unorganized (callus) tissues of the same genotype may provide an opportunity to address the problem of differential expression of genes regulating cytokinin metabolism during plant development.

[8-¹⁴C]Zeatin Metabolism in Phaseolus Embryos

by

Yun-Hwa Lee

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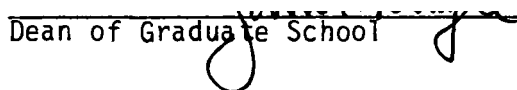
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LIST OF ABBREVIATIONS

Ade:	adenine
Ado:	adenosine
AMP:	adenosine-5'-monophosphate
ATP:	adenosine-5'-triphosphate
CI:	chemical ionization
DMSO:	dimethylsulfoxide
DZMP:	<u>trans</u> -dihydrozeatin 5'-monophosphate
DZ-O-G:	O-glucosyldihydrozeatin
EI:	electron impact ionization
GC-MS:	gas chromatography-mass spectrometry
HPLC:	High-performance Liquid Chromatography
i^6 Ado:	N^6 -(Δ^2 -isopentenyl) adenosine
i^6 Ade:	N^6 -(Δ^2 -isopentenyl) adenine
i^6 AMP:	N^6 -(Δ^2 -isopentenyl)adenosine 5'-monophosphate
ipn 6 Ade:	N^6 -(Δ^2 -isopentyl)adenine
ipn 6 Ado:	N^6 -(Δ^2 -isopentyl)adenine
RDZ:	dihydrozeatin riboside, 6-(3-methylbutylamino)-9-ribofuranosylpurine
RDZ-O-G:	O-glucosyl, 9-ribosyldihydrozeatin
RZ:	ribosylzeatin, 9- β -D-ribofuranosyl- <u>trans</u> -zeatin
<u>cis</u> -RZ:	<u>cis</u> -ribosylzeatin, 9- β -D-ribofuranosyl- <u>cis</u> -zeatin
RZ-O-G:	O-glucosyl, 9-ribosylzeatin
TBAP:	tetrabutylammonium phosphate
TEA:	triethylamine
TFA:	trifluoroacetic acid

[¹⁴C]Zeatin, [8-¹⁴C]Zeatin, Z: trans-zeatin, 6-(4-hydroxy-3-methyl-trans-2-butenyl-amino)purine

Z-7-G: 7-glucosylzeatin, 7-β-D-glucopyranosyl-trans-zeatin

Z-9-G: 9-glucosylzeatin, 9-β-D-glucopyranosyl-trans-zeatin

ZMP: trans-zeatin 5'-monophosphate

Z-0-G: 0-glucosylzeatin, 0-β-D-glucopyranosyl-trans-zeatin

[8-¹⁴C]Zeatin Metabolism in Phaseolus Embryos

I. INTRODUCTION

As part of a program to study the genetic regulation of cytokinin metabolism in Phaseolus, callus culture bioassays have been used to identify genotypic variations of interest. Genetic differences in cytokinin structure-activity relationships, cytokinin requirements (cytokinin-autonomous vs. cytokinin-dependent growth) and responses to phenylurea-type cytokinins have been detected (Mok et al., 1978; Mok et al., 1979; Mok et al., 1980; Mok et al., 1982a and Mok et al., 1982b). These studies have led to the characterization of interspecific differences in cytokinin destruction by nuclear genes and the identification of one major locus controlling cytokinin autonomy in P. vulgaris.

We are also interested in defining genetic variations in organized tissues of Phaseolus plants. Therefore, the metabolism of [¹⁴C]zeatin was examined in tissues of a number of Phaseolus species and genotypes. In the present paper we report the metabolism of [¹⁴C]-zeatin in immature embryos of P. vulgaris and P. lunatus.

II. LITERATURE REVIEW AND BACKGROUND INFORMATION

Cytokinins are compounds that promote cell division (Skoog et al., 1965). "Kinetin" was the first synthetic cytokinin (6-furfurylaminopurine) identified when Miller et al. (1955, 1956) isolated the substance upon dehydratation of DNA from herring sperm. Zeatin, found in immature corn kernels, was the first naturally occurring cytokinin to be isolated. Its structure was identified as 6-(4-hydroxy-3-methyl-trans-2-butenyl-amino)purine (Letham, 1963; Letham et al., 1964; Miller, 1961 and Letham and Miller, 1965) and was confirmed later by Shaw and Wilson (1964) via direct synthesis.

There are two broad classes of compounds that possess cytokinin activities (as defined by bioassays). The first group of compounds are N⁶-substituted amino purine analogues such as i⁶Ade, benzyladenine and zeatin. The second group are substituted phenylureas such as diphenylurea, N-phenyl-N'-(4-pyridyl)urea derivatives and N-phenyl-N'-1,2,3-thiadiazol-5-ylurea (thiadiazuran) (Shantz and Steward, 1955; Bruce and Zwar, 1966; Bruce et al., 1965; Isogai, 1981; Mok et al., 1976 and Mok et al., 1982a). Although the chemical structures of the two classes of compounds are distinct, the biological activities are similar. The mode of action of phenylurea type cytokinins in relation to that of adenine-type cytokinins has not been defined.

The structure-activity relationship of the compounds in the two classes have been studied, and were reviewed by Skoog and Armstrong

(1970), Strong (1958), Leonard (1974) and Matsubara (1980). Generally, an alkyl group as the N⁶-substituent (with the optimum length of 4-6 atoms) gives high activity. A double bond and a hydroxyl group further increase the activity. In addition, side chains which enhance the planarity of the molecule promote biological activity (Hecht et al., 1970). Among phenylurea-type cytokinins, substitutions in the phenylring, increase activity generally particularly with electronegative substituents.

A. Cytokinin Biosynthesis

Cytokinins occur naturally as components of specific t-RNA species, and as free forms in plant cells. At present, little is known about the mechanism of free cytokinin biosynthesis in plants. There are two hypotheses: degradation of t-RNA (Palni and Horgan, 1983) and de novo synthesis presumably via adenine or adenosine as precursors (Nishinari and Syono, 1980). Mevalonic acid (MVA) is generally considered as a precursor of the isoprenoid side chain. Both hypotheses have been reported but no definite evidence supporting either has been presented. Burrows and Fuell (1981) have suggested that free cytokinin production in cytokinin-autonomous and crown-gall tissues of tobacco takes place by a route not involving t-RNA. Palni and Horgan (1983) have found, however, the high level of trans-RZ observed from Vinca rosea crown-gall tissue is associated exclusively with the crown-gall tRNA. (For review see Letham and Palni, 1983).

B. Cytokinin Metabolism

Cytokinins have been found endogenously in a wide range of plant tissues. Naturally occurring cytokinins and some examples of the plant sources of these compounds are listed below: Z, from corn kernels (Letham, 1963) and cones of hops (Watanabe et al., 1981); cis-Z, from cones of hops (Watanabe et al., 1981); RZ, from corn kernels (Letham, 1966), Pinus radiata (Taylor et al., 1984) and cones of hops (Watanabe et al., 1981); cis-RZ, Z-7-G, from radish seeds (Summons et al., 1977); Z-9-G, from Vinca rosea crown-gall tissue (Peterson and Miller, 1977; Morris, 1977; Scott et al., 1980); RDZ and DZOG, from Phaseolus vulgaris leaves (Wang et al., 1977 and Wang and Horgan, 1978); Z-0-G, from Phaseolus vulgaris roots (Scott and Horgan, 1984a) and V. rosea crown-gall tissue (Peterson and Miller, 1977); RZOG, from V. rosea crown-gall tissue (Peterson and Miller; 1977) and cones of hops (Watanabe et al., 1981); cis-RZ, from cones of hops (Watanabe et al., 1981) and a glucoside of RZ in which the glucose moiety is attached directly to the ribose from Pinus radiata (Taylor, et al., 1984).

The metabolism of cytokinins has also been studied using radioactively labelled cytokinins applied exogenously to various plant organs, tissues, and cells.

Formation of the 7-glucoside of benzyladenine in tobacco callus was first demonstrated by Fox et al. (1973, 1974). Exogenously applied zeatin in radish roots and to derooted radish seedling (Parker and Letham, 1973; Cowley et al., 1978 and Gordon et al.,

1974) yield Ade, Ado, AMP, RZ, ZMP and Z-7-G. When [^3H] Zeatin was supplied to Zea mays L. seedlings (Cowley et al., 1978; Parker and Letham, 1974 and Parker et al., 1973) with roots excised, the metabolites were identified as AMP, Ado, Ade, Z-7-G (a minor metabolite). The principal metabolites formed from zeatin by the roots of intact Z. mays seedlings were Ade, Ado, AMP, RZ, ZMP, Z-7-G (a minor metabolite) and Z-9-G (a major metabolite). When zeatin was supplied to excised leaves of Populus alba, the principal metabolites formed were Ado, Z-0-G, DZ-0-G and RDZ-0-G; minor metabolites were AMP, Z-7-G, Z-9-G, DZ and RZ (Duke et al., 1979 and Letham et al., 1976). Labelled zeatin was supplied through the transpiration stream to derooted lupin (Lupinus angustifolius L.) seedlings. The major compounds were β - [6-(4-hydroxy-3-methylbut-trans-2-enylamino)-purin-9-yl]alanine (lupinic acid) and Z-0-G. The rest of the metabolites were DZ, RZ, RDZ, ZMP, DZMP, Z-7-G, Z-9-G, RZ-0-G, RDZ-0-G, dihydrolupinic acid, Ade, Ado and AMP (Parker et al., 1978; Duke et al., 1978 and Parker et al., 1975). All the above metabolites have been identified unequivocally by comparison with compounds obtained via direct syntheses.

The biological significance of various metabolites is still unclear. The O-glucosides and N-glucosides may be storage forms or may facilitate cytokinin transport (Letham et al., 1982 and Summons et al., 1980). They are also reported to be more stable and have higher resistance to some of the degradative enzymes.

C. Enzymes Involved in the Biosynthesis and Metabolism of Cytokinins

A number of enzymes involved in cytokinin biosynthesis and metabolism have been purified. Chen and Eckert (1977) reported the formation of $N^6-(\Delta^2\text{-isopentenyl})\text{adenosine-5'-monophosphate}$ ($i^6\text{AMP}$) from $i^6\text{Ado}$ and ATP by an adenosine kinase isolation from wheatgerm. In addition, nucleotides could be formed directly from the free base by the enzyme adenine phosphoribosyltransferase (Chen et al., 1982). A 5'-nucleotidase (Chen and Kristopeit, 1981a) converts nucleotides to corresponding nucleosides. The enzyme adenosine nucleosidase (Chen and Kristopeit, 1981b) catalyzes the deribosylation of $i^6\text{Ado}$ (to $i^6\text{Ade}$). Adenosine phosphorylase (Chen and Petschow, 1978) catalyzes the ribosylation of $i^6\text{Ade}$ to $i^6\text{Ado}$.

Paces et al., (1971) identified enzyme activity in crude extracts of tobacco tissues that could cleave the side chain of $i^6\text{Ado}$ and other adenine derivatives with unsaturated N^6 -side chains. This enzyme is similar to the cytokinin oxidase purified from Zea mays kernels by Witty and Hall (1974). The reaction converting $i^6\text{Ade}$, $i^6\text{Ado}$, Z and RZ to adenine and/or adenosine (McGaw and Horgan, 1983a) requires oxygen. The aldehyde, 3-methylbut-2-enal, is one of the intermediate products derived from the reaction of side chain cleavage (Brownlee et al., 1975). Dihydrozeatin (Palmer et al., 1981) and N^6 -benzyladenine (Parker et al., 1973) are not substrates for the enzyme. Cytokinin oxidase has also been purified from Vinca rosea tumor tissue, which is also similar to the maize enzyme referred to

above (Scott et al., 1982). Another cytokinin oxidase has been partially purified from callus tissues of Phaseolus vulgaris L. cv. Great Northern (Chatfield and Armstrong, 1984). This enzyme exhibits similar specificity for substrates, cytokinins with an unsaturated N⁶-side chain. The presence of glucosyl or ribosyl groups in the 7- or 9-position or an alanyl group in the 9-position of purine moiety has little effect on their susceptibility to cytokin oxidase, but O-glucosyl derivatives are resistant to oxidation (McGaw and Horgan, 1983b). In detached P. vulgaris leaves (Palmer et al., 1981) and V. rosea crown gall tissues (Horgan et al., 1981), side chain cleavage is also the fate of the majority of exogenously applied zeatin.

Cytokinin can occur naturally as glucosides with β -D-glucose as the sugar substituent. An enzyme has been purified from radish (Raphanus satives) cotyledons, cytokinin-7-glucosyltransferase, which utilizes uridine diphosphate glucose as the glucose donor and converts zeatin into its 7- and 9-glucosides (Entsch et al., 1979).

β -(9-Cytokinin)alanine synthase, derived from immature Lupinus luteus seeds, required the unusual and unstable substrate O-acetylserine as donor of the alanine moiety and converted zeatin to 9-alanyl-zeatin (Murakoski et al., 1977 and Entsch et al., 1983). Such an enzyme may also be classified as a C-N-ligase (Entsch et al., 1983).

D. Analysis of Cytokinins

Cytokinins and their metabolites are usually extracted with

organic solvent or acids and purified and analyzed by chromatography. Thin layer paper and liquid chromatography have been used extensively to separate and identify cytokinins.

Carnes et al. (1975) first introduced HPLC for the fractionation and analysis of cytokinins. The method is suitable in separating cytokinin bases, nucleosides, glucosides and nucleotides (Horgan and Kramers, 1979 and Scott and Horgan, 1982). The cis- and trans-forms of zeatin and ribosylzeatin can be separated by increasing the concentrations of the organic phase. The analysis time using HPLC is shorter and the resolution is much higher than low pressure column chromatography such as polystyrene-base ion exchange resins and Sephadex LH-20 columns. The final HPLC fractions are often pure enough for direct GC-MS analysis.

Mass spectrometry has been used to determine the molecular weights, mass ion composition and quantity of cytokinins. Generally there are two ways of converting the purified metabolites to volatile derivatives via methylation: trimethylation (TMS) and permethylation. Some metabolites are more amenable to one or the other method of methylation. For example: MacLeod et al. (1976) compared the suitability of TMS and permethylated derivatives of glucosides of zeatin and N⁶-benzyladenine for GC-MS and mass spectral studies. Mass spectra of the TMS derivatives show more significant isomeric differences than the corresponding permethylated compounds and this method of derivatization also gave better results for N-glucosides. However, the permethylation has the advantage of being hydrolytically

stable and the derivatized products are lower in molecular weight than the TMS derivatives. Scott and Horgan (1984b) recently compared the mass spectra of cytokinin metabolites from tobacco crown gall tissue using both derivatizations.

Both electron impact (EI) ionization and chemical ionization (CI) mass spectrometry can be used for the identification of purified metabolites. The CI technique is useful for cytokinins having a sugar moiety linked through the side chain oxygen atom (Summons et al., 1980). But CI-MS gives less fragmentations compared with EI-MS. An ideal approach would be using both EI and CI spectro-analyses. Usually, direct synthesis is performed to unequivocally confirm the structure of new metabolites identified.

E. Cytokinin Metabolism in Phaseolus Species

Studies of cytokinin metabolism in intact plant tissue of Phaseolus species have been examined (Wang and Horgan, 1978; Palmer et al., 1981 and Scott and Horgan, 1984a). The major cytokinin identified in P. vulgaris leaves and decapitated plants was DZ-0-G and the minor cytokinin was RDZ (Wang et al., 1977 and Wang and Horgan, 1978). The quantity of DZ-0-G was determined using the isotope dilution method employing a deuterium-labelled internal standard and combined GC-MS (Palmer et al., 1981). The accumulation of cytokinin appeared to parallel the gradual increase in fresh weight of leaves of intact plants, or in leaves of plants that were decapitated but not disbudded. When secondary lateral buds were allowed to grow out

from decapitated plants the levels of DZ-0-G in the primary leaves rapidly declined to a value similar to or lower than that found in leaves of intact plants. The major cytokinins in stems of decapitated, disbudded bean plants have been identified as RZ, RDZ, ZMP and DZMP. The stability of cytokinin has been examined (Palmer, et al., 1981). The order of the stability appears to be DZ-0-G > Z-0-G > DZ > Z. These results suggest that side-chain glucosylation may confer some degree of metabolic stability on cytokinins in bean leaves although it is not clear whether this effect might be due to an inhibition of enzymic cleavage of the side-chain or to compartmentation of the glucoside away from the site of metabolism.

Callus cultures have been used to study the genetic variations in cytokinin metabolism. Studies on the structure-activity relationships of tissue cultures derived from P. vulgaris cv. Great Northern and P. lunatus cv. Kingston revealed dramatic differences in the responses of these callus tissues to cytokinins bearing unsaturated isoprenoid side chain [zeatin and $i^6\text{Ado}$] (Mok et al., 1978). In P. vulgaris cultures, the presence of a double bond in the cytokinin side chain resulted in a marked reduction in cytokinin activity. The growth responses of callus of the interspecific hybrid were intermediate between the parental tissue (Mok et al., 1982b). The differential structure-activity relationship was related to a rapid degradation of unsaturated cytokinins in P. vulgaris tissue (Mok et al., 1982b), presumably due to a higher level of cytokinin oxidase.

In order to determine if genetic variations in cytokinin metabolism also occur in whole plant tissues of Phaseolus, [^{14}C]zeatin metabolism was examined in a variety of tissues. The research described in this dissertation concerns [^{14}C]zeatin metabolism in immature embryos of P. vulgaris and P. lunatus. The results are presented in manuscript format.

III. MATERIALS AND METHODS

Plant Materials

Seeds of Phaseolus vulgaris L. cv. Great Northern (GN) and P. lunatus cv. Kingston (K) were originally obtained from Dr. Dermot Coyne (University of Nebraska) and Dr. Don Grabe (Oregon State University). Plants were grown in the greenhouse at 25°C with the photoperiod of 14 h. Immature embryos, 3, 6, and 9 mm in length were collected.

Chemicals

Trans-zeatin, trans-ribosylzeatin, Ade, Ado, 5'-nucleotidase (Crotalus adamanteus venom) and β -glucosidase (almond) were obtained from Sigma. 9-Glucosylzeatin was a gift from Dr. R. Durley (Oregon State University). 6-Chloro [8- ^{14}C]purine (24 mCi/mmol) was obtained from Amersham. [8- ^{14}C]Zeatin (24 mCi/mmol) was synthesized from 6-chloro-[8- ^{14}C]purine following procedures published elsewhere. (Kadair et al., 1984).

Metabolism of [^{14}C]zeatin

Immature embryos at three developmental stages (measuring 3, 6, and 9 mm in length) were dissected from the pods under sterial condition. The embryos corresponded to late heart, and early and mid

cotyledonary stages at the respective length. [^{14}C]Zeatin (0.05 μCi , 0.002 μmol) dissolved in 250 μl of H_2O was applied aseptically to 250 mg of immature embryos. The vials were sealed and maintained at 27°C in the dark for 2, 4, and 8 hours. In addition, embryos were incubated with radioactively labeled zeatin dissolved in 0.05 M Tris-HCl (pH 6.0) buffer with all other conditions identical to those described above. To determine the amount of radioactivity recovered at time 0, [^{14}C] zeatin was applied and metabolites extracted immediately. This estimate was taken for the three sizes of embryos of both genotypes. Each experiment was repeated at least once and averages of two experiments are presented.

To extract metabolites, the embryos were homogenized with a Tissuemizer equipped with a Microprobe Shaft (Tekmar) in 2ml of cold 95% ethanol. Cell debris was removed by successive filtration through Whatman paper (No. 1) and Millipore filters (0.25 μm). The ethanol extract was taken to dryness in vacuo at 35°C , redissolved in 2 ml of 50% (v/v) ethanol and centrifuged at 23,500 g for 20 min. The supernatant was condensed in vacuo to 100 μl with a speed vac concentrator (Savant). The sample was then analysed directly by HPLC. A Beckman Model 110 dual pump HPLC with a prepacked column of reversed-phase C_{18} (Ultrasphere ODS 5 μm , 4.6 x 250 mm; Altex) was used. The aqueous buffer consisted of 0.2 M acetic acid, adjusted to pH 3.5 with TEA. Samples were eluted with a linear gradient of methanol (5-50% over 90 min) in TEA buffer at a flow rate of 1 ml/min. Fractions of 1 ml were collected and counted in Ready-Solve

MP Scintillation fluid (Beckman) with a Beckman LS 7000 Scintillation counter. To purify and further analyse nucleotides, paired-ion reversed-phase HPLC (Capelle et al., 1983) was used. The buffer consisted of 0.3% (w/v) TBAP and 0.65% (w/v) KH_2PO_4 , adjusted to pH 5.8 with NH_4OH , and acetonitrile was used as the organic phase. The sample was applied in 100 μl of 20% acetonitrile in buffer and eluted with a linear gradient of acetonitrile (20-60% over 20 min) in buffer at a flow rate of 1.5 ml/min. Fractions of 1 ml were collected and counted as described above. To purify other metabolites, a reversed-phase C_{18} column was used but with a buffer solutions of 0.2 M acetic acid adjusted to pH 4.8 with TEA. The metabolites were eluted by a methanol gradient of 5-50% over 90 min at a flow rate of 1 ml/min.

Uptake of [^{14}C]zeatin and distribution of metabolites
in embryos and solution

To determine if there are differences in the uptake of [^{14}C]zeatin between GN and K embryos and the distribution of metabolites in embryonic tissues vs. incubating solution, the following experiments were performed. [^{14}C]Zeatin (0.1 μCi ; 0.004 μmol) dissolved in 250 μl of H_2O was incubated with 250 mg of immature embryos. (The amount of radioactively labeled zeatin was twice that of other experiments in order to obtain sufficient radioactivity in HPLC fractions when embryo and solution portions are analyzed separately). After 2 h, the embryos were removed and subsequently washed with 1 μM of cold zeatin in 1.5 ml of H_2O . The wash solution

was combined with the incubating solution, dried in vacuo and analysed with reversed-phase HPLC. The embryos were homogenized and zeatin metabolites were extracted and analyzed as described above. The same experiment was performed for embryos of each size class.

Identification of [¹⁴C]zeatin metabolites

Treatment with 5'-nucleotidase: Fractions collected after HPLC analysis were dried and redissolved in 0.05 M Tris-HCl buffer (pH 6.8) containing 5 mM MgCl₂, and aliquots of 30 μ l were incubated with one unit of 5'-nucleotidase or 3'-nucleotidase for 0.5 h at 37 °C (Aung, 1978). Ethanol (1.5 ml) was added and the solution was centrifuged at 23,500 g for 20 min. The supernatant was taken to dryness in vacuo at room temperature, redissolved in 100 μ l of 5% methanol and fractionated by HPLC.

Treatment with β -glucosidase: Fractions collected after HPLC analysis were dried and redissolved in 200 μ l of 0.03 M acetate buffer (pH 5.3). After adding 0.5 units of the enzyme, the solution was incubated at 37°C for 1 h. Ethanol (95%, 1.5 ml) was added and the solution was centrifuged at 23,500 g for 20 min. The supernatant was taken to dryness in vacuo at room temperature, redissolved in 100 μ l of 5% methanol and analysed by HPLC.

Treatment with periodate and cyclohexamide: Fractions collected after HPLC analysis were dried and dissolved in 10 μ l of periodate (10% w/v). After 3 h at room temperature, 2 μ l of cyclohexamide was

added (Robins et al., 1967). After 18 h the solution was dried and the residue redissolved in 100 μ l of 5 % methanol and analysed by HPLC.

Treatment with permanganate: Fractions collected after HPLC analysis were dried and redissolved in 500 μ l of H_2O , and 250 μ l of 0.1 N $KMnO_4$ (w/v) was added (Aung, 1978 and Hall, 1971). After 5 min, 1 ml of 95% ethanol was added and the solution was left for 24 h at room temperature. After centrifugation at 23,500 g, the supernatant was dried in vacuo. The residue was redissolved in 100 μ l of 5% methanol and analysed by HPLC.

Acid hydrolysis with TFA: Fractions collected from HPLC analysis were dried and redissolved in 1 ml of 0.6 M TFA. The solution was incubated at 95°C for 3 h (Hall, 1964; Letham et al., 1979 and Miura and Hall, 1973) and centrifuged after addition of 1 ml H_2O . The supernatant was dried in vacuo, redissolved in 5% methanol and chromatographed by HPLC.

Permethylation of metabolites for GC-MS analysis: Anhydrous DMSO was obtained by treating DMSO with activated molecular sieves over night. Methyl sulfinyl anion (4% w/v) was prepared by reacting NaH (200 mg) with 5 ml of DMSO under nitrogen. Standards and/or metabolites were dried and dissolved in 100 μ l of anhydrous DMSO, followed by addition of 25 μ l of methyl sulfinyl anion solution after 15 minutes by 10 μ l of CH_3I . After 1.5 h, the reaction was terminated by adding 1 ml of H_2O . Permethylated derivatives were extracted with 1 ml of $CHCl_3$. The organic layer was washed three

times with 1 ml H₂O, dried and redissolved in 5 ul of CHCl₃ prior to GC-MS analysis.

GC-MS analysis: GC-MS analyses were performed using a Finnigan model 4023 GC-MS computer system with model 4500 source retrofit and pulsed positive-negative chemical ionization. The source temperature was 190°C. The conditions for EI and CI were 70 eV electron energy and 0.75t methane pressure in the source with 70 eV electron energy and 0.75t methane pressure in the source with 70 eV electrons respectively. For gas chromatography, a glass column (250 mm x 2 mm ID, Pyrex) packed with 7% OV-101 on Supelcoport (100-120 mesh) was used. The column temperature was programmed to run from 200 to 320°C (8°C/min). The injector and detector temperature was 275°C. The capillary (15 m x 0.25 mm ID) contained fused silica (J and W DB-1). For combined GC-MS analyses, the temperature was held at 50°C for 1 min, raised to 225°C (25°C/min) and then to 320°C (8°C/min).

IV RESULTS

[¹⁴C] Zeatin metabolism in Phaseolus vulgaris L. cv. GN. and
P. lunatus cv. K.

Ethanol extracts of embryos of P. vulgaris (GN) and P. lunatus (K) were fractionated by HPLC. Representative elution profiles (3 mm embryos, 4 h incubations) are presented in Figs. 1 and 2 respectively for the two genotypes. In both genotypes [¹⁴C]zeatin was rapidly metabolized. Five major radioactive peaks in addition to zeatin were recovered in extracts of GN (Fig. 1). One of the peaks coeluted with ribosylzeatin (see Fig. 3 for the elution positions of standards); the remaining four peaks were designated as A (35-36), I (45-47), II (58-59) and C (62-63). Large amounts of compound I were formed in embryos of this genotype. In K extracts, however, metabolites I and II were absent (Fig. 2). Of the four major metabolites recovered from this genotype, three had elution positions identical to those of ribosylzeatin, A and C. However, the fourth metabolite (fractions 37-38) was not found in GN extracts and was designated as B.

Identification of major radioactive metabolites:

Metabolite A:

Metabolite A from both tissues was treated with 5'-nucleotidase and rechromatographed on HPLC. The radioactivity co-eluted with ribosylzeatin (Fig. 4A). Treatment with 3'-nucleotidase had no effect on metabolite A. These results suggest that A is a 5'-

FIGURE 1. HPLC profiles of [^{14}C] zeatin metabolites obtained from embryo extracts (3 mm, 4 h incubation) of P. vulgaris cv. GN.

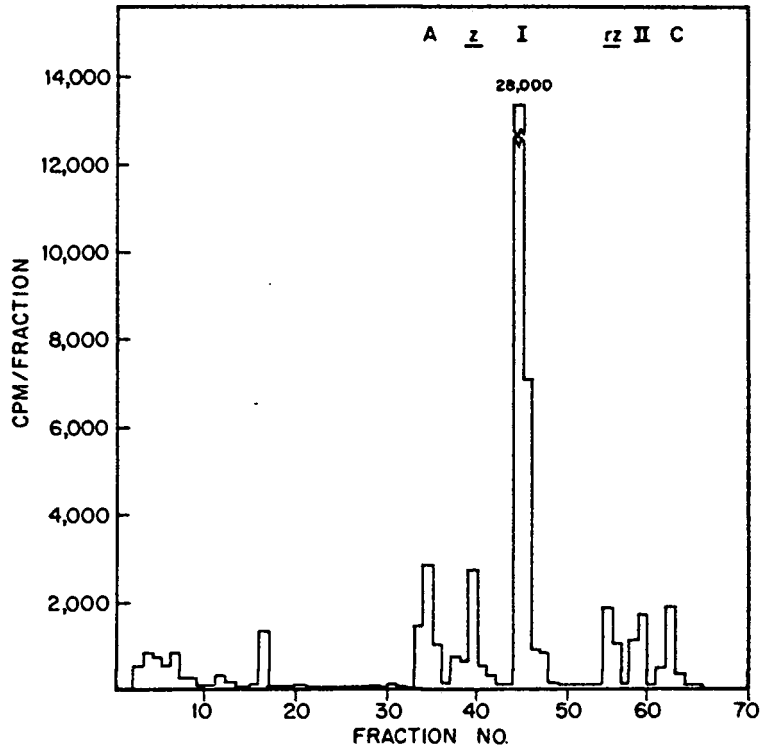


Figure 1

FIGURE 2. HPLC profiles of [^{14}C]zeatin metabolites obtained from embryo extracts (3 mm, 4 h incubation) of P. lunatus cv. K.

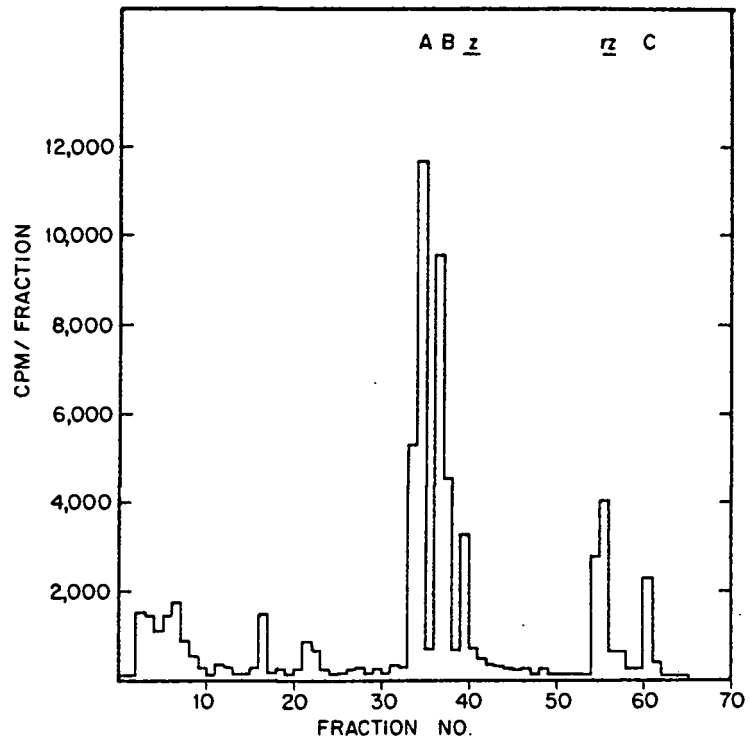


Figure 2.

FIGURE 3. HPLC profiles of Ade, Ado and zeatin standards.

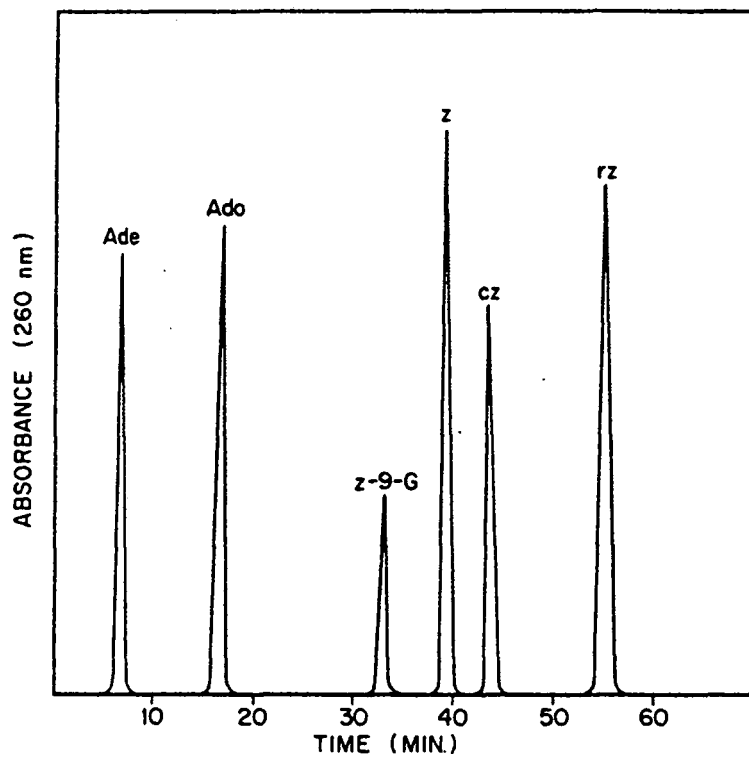


Figure 3.

FIGURE 4. Analyses by HPLC of metabolites A after treatment with 5'-nucleotidase (A), B after treatment with β -glucosidase (B), and C after treatment with β -glucosidase (C). Bars with discontinuous border indicate elution position of radioactivity before treatments.

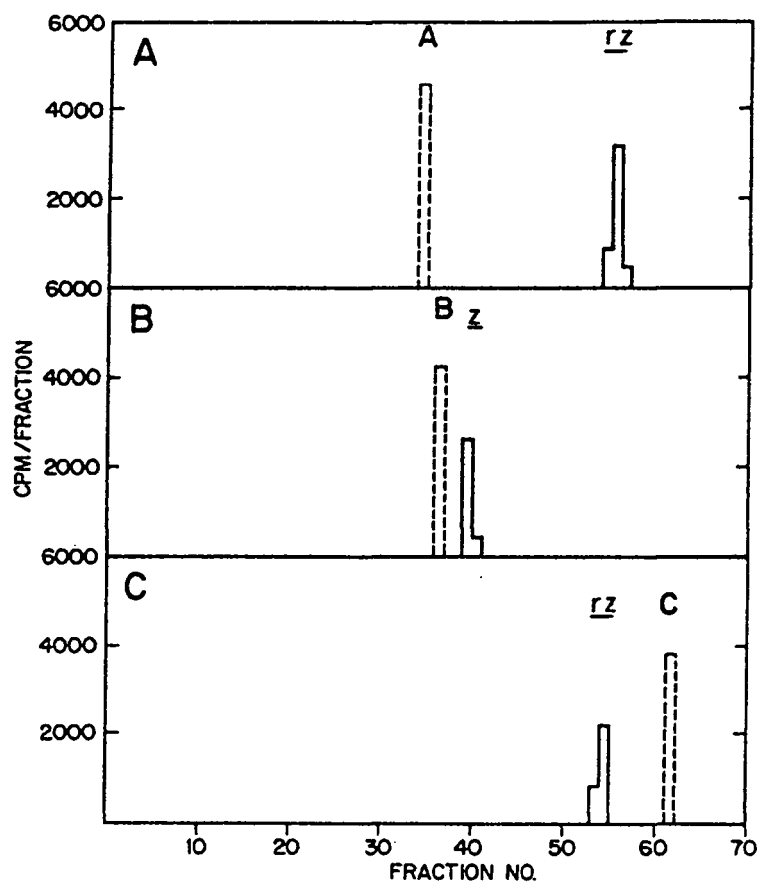


Figure 4.

nucleotide of zeatin. To determine if A is the mono-, di- or trinucleotide, it was treated with KMnO_4 and rechromatographed on paired-ion reversed-phase HPLC (Capelle et al., 1983). The radioactivity co-eluted with AMP. Thus metabolite A seems to be the 5'-mononucleotide of zeatin.

Metabolite B:

Metabolite B from tissue extracts was treated with β -glucosidase. After fractionation by HPLC the radioactivity coincided with the elution position of zeatin (Fig. 4B). Thus this metabolite seems to be the O-glucoside of zeatin.

Metabolite C:

Treatment of metabolite C with β -glucosidase resulted in a shift of the elution position to that of ribosylzeatin after HPLC fractionation (Fig. 4C). Acid hydrolysis with TFA (removing sugar moieties (Hall, 1964; Kadair et al., 1984 and Miura and Hall, 1973)) shifted the radioactivity to the position of zeatin. Therefore, metabolite C is most likely the O-glucoside of ribosylzeatin (Entsch et al., 1980 and Horgan, 1975). Similar results were obtained for metabolite C recovered from the two genotypes.

Metabolites I and II:

Metabolites I and II were treated with β -glucosidase. About one-half of I was converted to zeatin (Fig. 5A), while one-half of the

FIGURE 5. Analyses by HPLC of metabolite I after treatments with β -glucosidase (A), TFA (B), periodate plus cyclohexamide (C) and permanganate (D). Bars with discontinuous border indicate elution position of radioactivity before treatments.

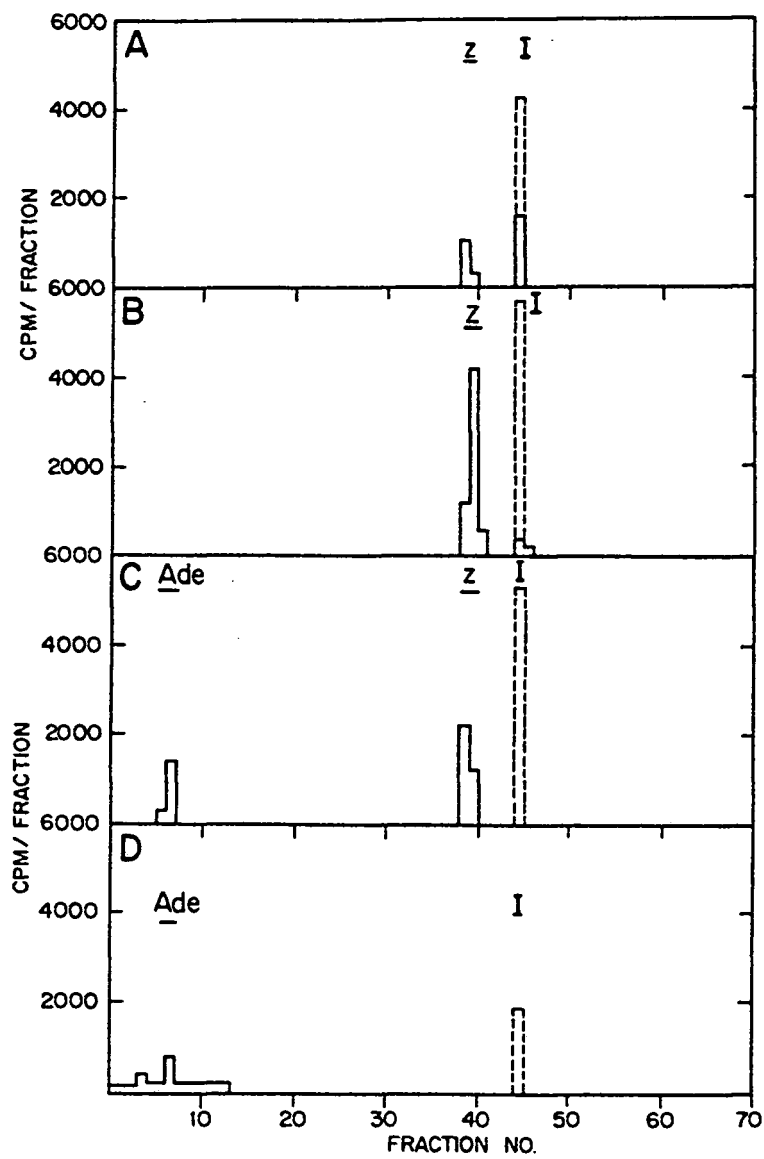


Figure 5.

original radioactivity of II shifted to the position of ribosylzeatin (Fig. 6A). These results suggest that portions of metabolites I and II probably involve glycosylation of zeatin and ribosylzeatin. The difference between I and II may reside in the ribosylation of II at the 9 position. Acid hydrolysis with TFA converted the majority of I to zeatin (Fig. 5B), whereas identical treatment of II resulted in the formation of zeatin and small radioactive peak eluting off at the position of I (Fig. 6B). Treating metabolites I and II with periodate and cyclohexamide resulted in the recovery of zeatin in both cases (Figs. 5C and 6C). (Periodate opens ring structures of sugar moieties at the 2-3 position allowing subsequent removal of the open ring by cyclohexamine (Robins *et al.*, 1967)). Finally, metabolites I and II were treated with KMnO_4 to remove the N^6 -sidechain (Aung, 1978 and Hall, 1971). Compound I was converted to Ade and other breakdown products (Fig. 5D) indicating that the structure modification resides on the N^6 -sidechain. The Majority of metabolite II was converted to Ado and Ade (Fig. 6d). Based on the results of chemical and enzymatic tests, it is reasonable to conclude that metabolites I and II are most likely O-glycosylated (by not O-glucosylated) derivatives of zeatin and ribosylzeatin respectively.

To further test our interpretation, metabolite I was re-purified by HPLC with paired-ion reversed-phase (pH 5.8) and with reversed-phase C_{18} (pH 4.8) columns. The UV spectra of the purified compound at acidic (pH 3.5) and alkaline (pH 8.5) conditions are presented in Fig. 7. The absorption maxima were at 272 and 270 nm for the two

FIGURE 6. Analyses by HPLC of metabolite II after treatments with β -glucosidase (Fig. A), TFA (Fig. B), periodate plus cyclohexamide (Fig. C) and permanganate (Fig. D). Bars with discontinuous border indicate elution position of radioactivity before treatments.

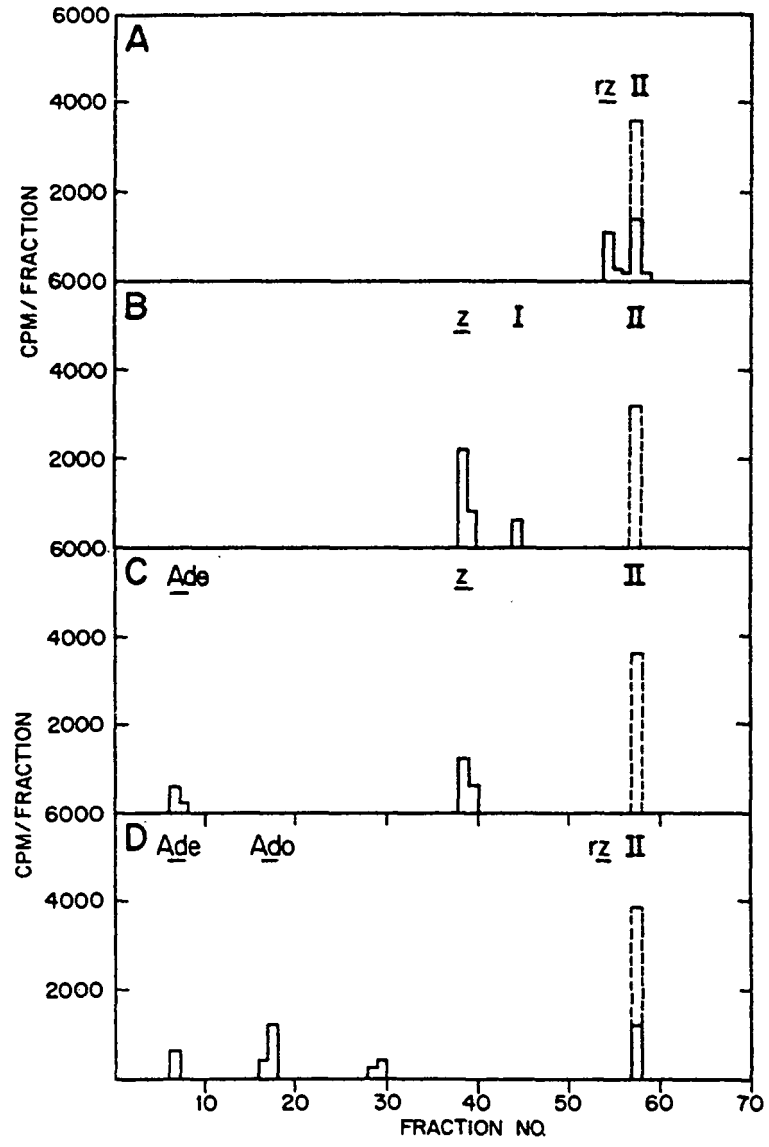


Figure 6.

FIGURE 7. UV spectra of purified metabolite I at pH 3.5 (A) and pH 8.5 (B).

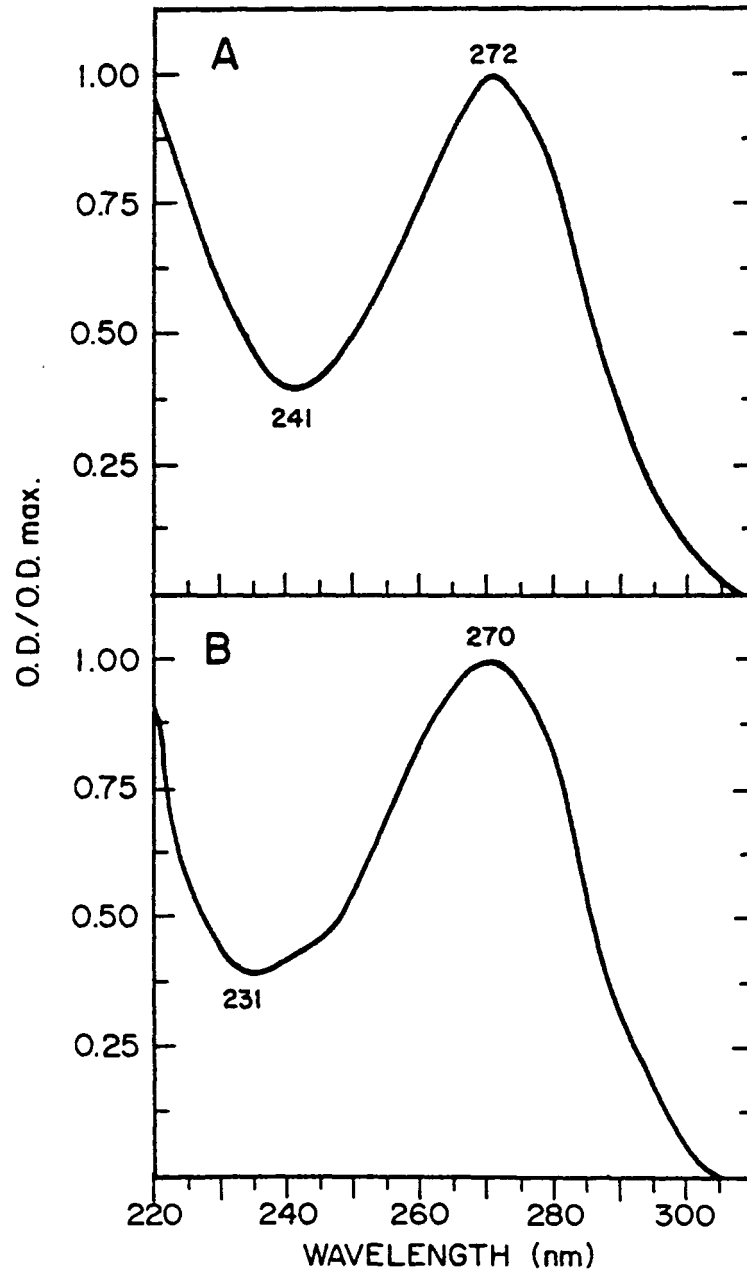


Figure 7.

conditions. The permethylated derivative of I was prepared and subjected to GC-MS analyses (Fig. 8). The mass spectrum obtained by CI (Fig. 8C) indicates a molecular weight of 421, identical to that of ribosylzeatin (Fig. 8B). The fragmentation pattern of lower molecular weight ions was similar to that of zeatin (Fig. 8A). As mass spectral analysis by EI has the advantage of generating larger number of molecular fragments, permethylated zeatin, ribosylzeatin and compound I derivatives were also subjected to this type of MS analysis (Figs. 8D, E and F). The spectrum obtained confirmed the molecular weight of 421 for metabolite I. The fragmentation pattern (Fig. 8F) was consistent with our interpretation that compound I could be an O-glycoside of zeatin. As the molecular weight indicates the presence of a pentose sugar, compound I is most likely O-ribosylzeatin.

Since both metabolites I and II had the tendency of premature fragmentation under EI, much larger amounts of sample (approximately 2 ug) were required to obtain a complete spectrum. The relatively small amount of metabolite II obtained dictated an analysis by CI only (Fig 9). The GC-MS analyses revealed a molecular weight of 581. The pattern of fragmentation was compatible with the structure of the O-riboside of ribosylzeatin. Nevertheless, direct synthesis of these compounds is needed to unequivocally confirm our interpretation of the structures.

FIGURE 8. Mass spectra of permethylated zeatin (A and D), ribosylzeatin (B and E) and metabolite I (C and F) obtained by CI (A, B and C) and EI (D, E and F) analyses.

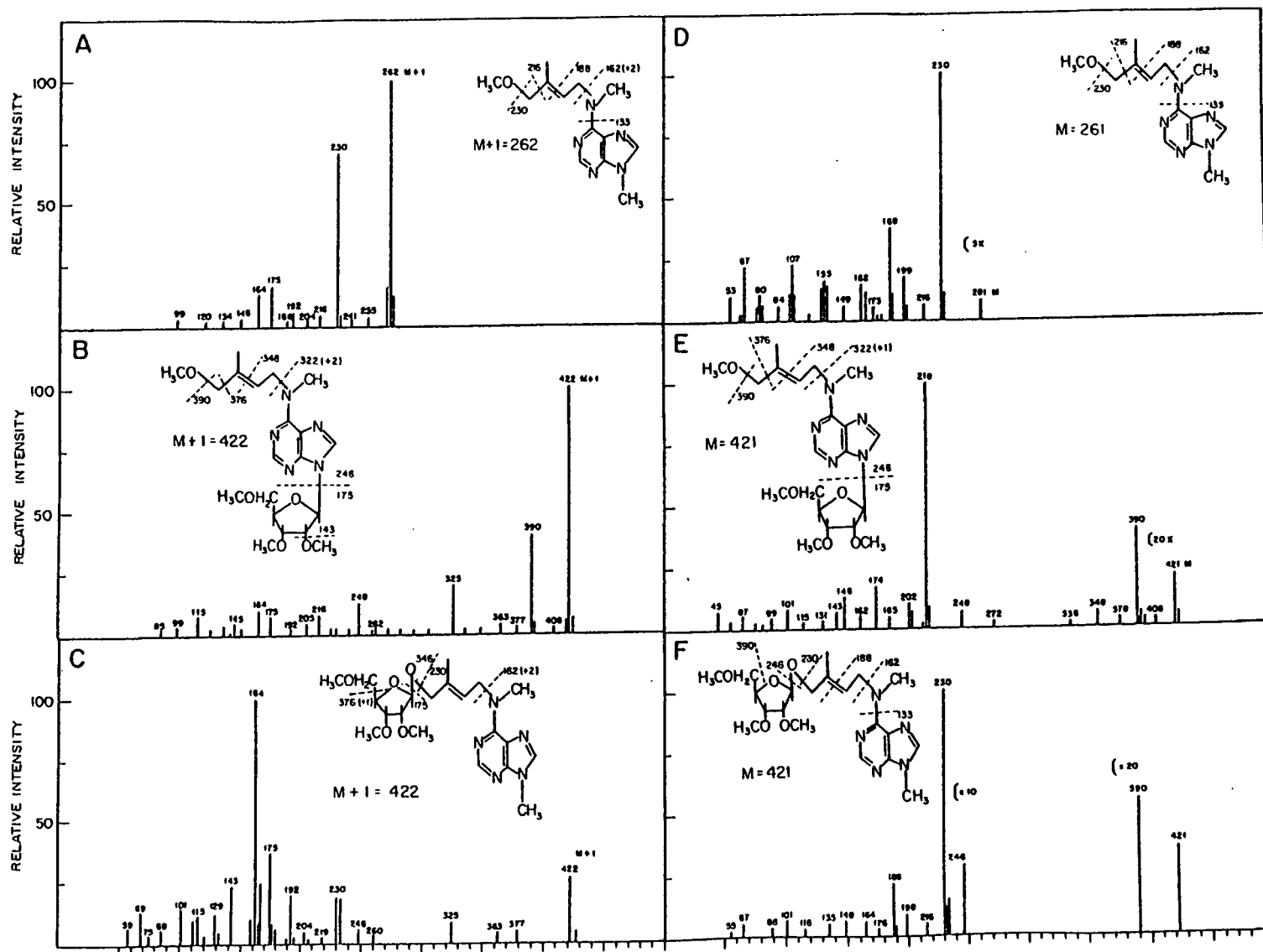


Figure 8.

FIGURE 9. Mass spectrum of permethylated metabolite II obtained by CI analysis.

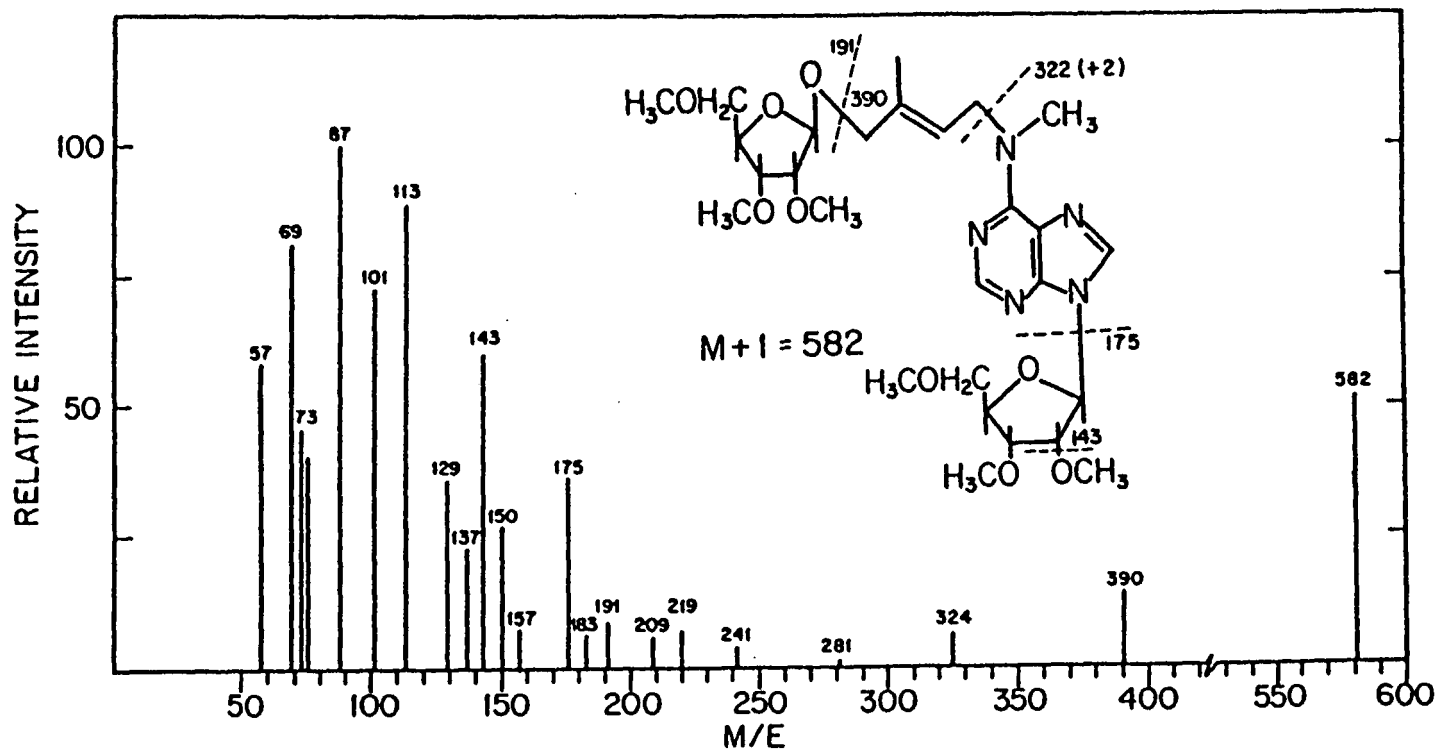


Figure 9.

Effects of embryo size on the metabolism of [^{14}C]zeatin

The proportion of radioactive metabolites (presented as % radioactivity at time 0) recovered from different sizes of embryos at various incubation times (2, 4 and 8 h) are presented in Fig. 10. No qualitative differences were observed between the 3, 6 and 9 mm embryos. However, some differences in the quantities of metabolites were observed between developmental stages. Younger embryos (3 and 6 mm) of GN appeared to convert larger proportions of zeatin to compounds I and II. Moreover, generally higher levels of zeatin mononucleotide were recovered from K embryos than GN embryos at all developmental stages. After 8 h, little radioactivity remained.

Distribution of metabolites in embryonic tissues vs. incubating solution

The distribution of metabolites in embryonic tissues vs. incubating solution (% of radioactivity recovered) is presented in Fig 11. Most of the zeatin metabolites were recovered from the embryos as well as the incubating solution, indicating that most likely the metabolites (and possibly also some of the enzymes) are released from the embryos to the solution. However, the majority of the zeatin nucleotide was retained in the embryonic tissues which is consistent with the reported properties of nucleotides. Also ribosylzeatin-0-glucoside was predominately found in the embryonic tissues, in contrast to zeatin-0-glucoside which was present mainly in the solution.

FIGURE 10. Radioactive metabolites recovered from embryo extracts of P. vulgaris cv. GN and P. lunatus cv. K.

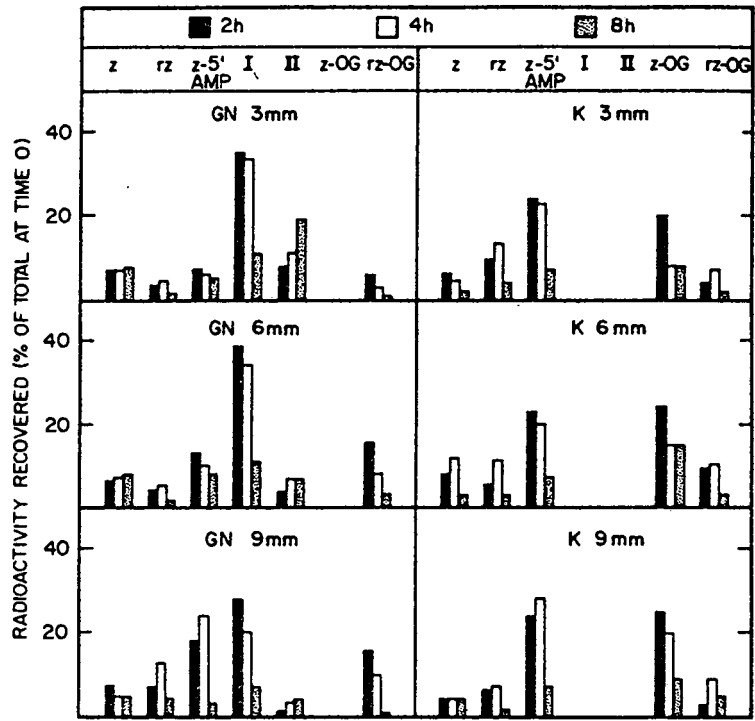


Figure 10.

FIGURE 11. Distribution of [^{14}C]zeatin metabolite between embryonic tissues and incubating solution.

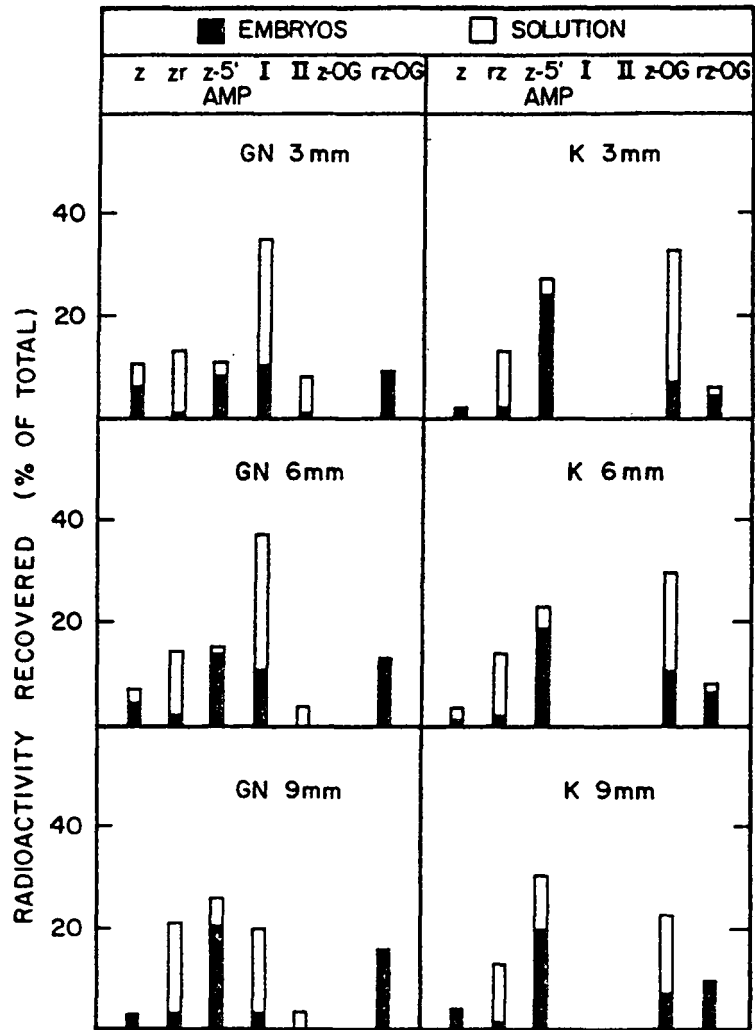


Figure 11.

Incubating embryos with radioactive zeatin dissolved in water and Tris-HCl buffer (in selected samples) gave essentially the same results regarding the types and proportion of metabolites recovered as well as the distribution of metabolites between embryonic tissues and incubating solutions.

V. DISCUSSION

The most striking difference in zeatin metabolism between P. vulgaris and P. lunatus embryos resides in the occurrence of metabolites I and II in P. vulgaris only. These compounds were tentatively identified as the O-ribosides of zeatin and ribosylzeatin. As far as we know, the ribosylation of the N⁶-sidechain of zeatin has not been previously reported (Letham and Palni, 1983). Since earlier studies of zeatin metabolism in P. vulgaris axes, seeds, leaves and roots (Palmer et al., 1981; Sondheimer and Tzou, 1971 and Wang et al., 1977) yielded primarily zeatin glucosides, nucleotides, and dihydrozeatin derivatives, the occurrence of metabolites I and II appears to be restricted to embryonic tissues. Unpublished results obtained in our laboratories using other plant parts of cv. GN also substantiate the embryo-specific nature of these compounds. In addition, their presence is also likely to be species-specific, since two additional genotypes of P. vulgaris and P. lunatus have been examined and the results obtained were similar to those reported in this study for GN and K respectively.

The biological significance of O-ribosylation is presently unknown. However, the conversion of close to 30% of the exogenously supplied zeatin to compounds I and II in relatively short periods of time must represent a rather active process. Moreover, the enzyme system(s) associated with the ribosylation of the sidechain must be

quite specific since glucosylation of the sidechain and the ribosylation at the 9 position of the purine ring occur in GN as well as K. Therefore, intuitively, one would tend to speculate that the occurrence of these metabolites may be important to some aspects of embryonic growth in P. vulgaris. The immature embryos used in this study correspond to late heart (3 mm), early and mid-cotyledonary (6 and 9 mm) stages, with the younger embryos undergoing rapid growth (Walbot et al., 1972). It is of interest to note that younger embryos were also more active in the formation of metabolite I.

The biological functions of zeatin metabolites (such as O- and N-glucosides (Fox et al., 1974; Horgan, 1975 and Parker et al., 1973), lupinic acid (Letham et al., 1979; Murakoshi et al., 1977 and Parker et al., 1978), glucosyl-ribosylzeatin (Taylor et al., 1984), nucleoside and nucleotides (Laloue et al., 1974 and Sondheimer and Tzou, 1971)) are not well established. Glucosides have been suggested to be either storage forms or to be related to cytokinin transport (Gordon et al., 1974 and Letham and Palni, 1983). Protection against cytokinin oxidase (Whitty and Hall, 1974) attack via O-glucosylation and ribonucleotide formation have also been postulated (McGaw and Horgan, 1983b). Zeatin-9-riboside could be an intermediate in nucleotide formation or breakdown of nucleic acids. The occurrence of O-ribosylation of zeatin, however, would seem to suggest additional roles of ribosylation of cytokinin bases.

The development of interspecific hybrid embryos of Phaseolus has previously been examined in some detail (Mok et al., 1978 and

Rabakoarihanta et al., 1979). The arrest of embryo growth is dependent on the species combination and the direction of the cross. For example, P. vulgaris x P. lunatus embryos develop only to the preheart stage. The reciprocal cross gives embryos which cease to divide at the four-celled stage, but can be stimulated to progress to the preheart stage by supplying cytokinin to the female parent via hydroponic culture (Mok et al., in press). P. vulgaris x P. acutifolius crosses result in embryos which develop to the cotyledonary stage but also do not reach maturity. It was reported that the slower growth rate of P. vulgaris-P. acutifolius hybrid embryos was correlated with lower amount of extractable cytokinis as compared with selfed embryos of both species (Nesling and Morris, 1979). As the largest differences in cytokinin metabolism usually occur between Phaseolus species, it is conceivable that an imbalance of cytokinin utilization or metabolism in the hybrid tissues could have contributed to the abnormal growth of the interspecific hybrid embryos. The qualitative difference in zeatin metabolism between P. vulgaris and P. lunatus embryos reported here provides a useful basis for further research into the relationship between abnormal embryonic development and hormonal derangement.

Identification of genetic variations is essential for studies of the genetic regulation of metabolic processes. Callus cultures have proven to be a versatile screening system for genetic variations in cytokinin metabolism. By utilizing cell culture systems, it has been discovered that there is a differential structure-activity relation-

ship between P. vulgaris and P. lunatus (Mok et al., 1978). The substantially lower activity of cytokinins with an unsaturated N⁶-sidechain (zeatin, i⁶Ade) in the P. vulgaris callus bioassay was found to be related to a rapid degradation of the N⁶-sidechain of i⁶Ado (Mok et al., 1982). Another genetic variation detected in cell cultures was cytokinin autonomous growth of P. vulgaris tissues which was controlled by a major locus (Mok et al., 1980). The results described in this paper indicate that intrinsic genetic differences can also be detected at the whole plant level. Since these differences are qualitative and tissue-specific, analyses of both organized (intact) and unorganized (callus) tissues of the same genotype may provide an opportunity to address the problem of differential expression during development of the genes governing cytokinin metabolism.

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