# Biosynthesis of Cyclopiazonic Acids in Penicillium cyclopium: The Isolation of Dimethylallylpyrophosphate:cyclo-Acetoacetyltryptophanyl Dimethylallyltransferase

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During the production of  $\beta$ -cyclopiazonic acid ( $\beta$ CA) by *Penicillium cyclopium*, the following points are noted: (a) Dimethylallylpyrophosphate (DMAPP), which is incorporated into  $\alpha$ -cyclopiazonic acid ( $\alpha$ CA) in vivo, stimulates overall CA synthesis, whereas tryptophan, although incorporated into  $\alpha$ CA, inhibits overall CA synthesis. (b) A previously suggested substrate,  $\gamma$ ,  $\gamma$ -dimethylallyltryptophan, is not a precursor of  $\alpha$ CA. (c) The accumulation of *cyclo*-acetoacetyl-L-tryptophanyl (cAATrp) is described in both the culture medium and mycelium with increasing growth. (d) A cell-free extract of mycelium will catalyze the conversion of exogenous cAATrp and exogenous DMAPP into  $\beta$ CA in 1:1 stoichiometry, the  $\beta$ CA being bound to a protein.

## INTRODUCTION

In 1968 Holzapfel (1) isolated and characterized cyclopiazonic acid  $(\alpha CA)$ , a toxin from *Penicillium cyclopium* Westling strain 1082. Later Holzapfel et al. (2) discovered a new metabolite, *bissecodehydrocyclopiazonic* acid  $(\beta CA)$ , which was shown to be a direct precursor of  $\alpha CA$ . This was later confirmed by the isolation of five isoenzymes which caused the transformation (3). During the conversion of  $\beta$ - to  $\alpha CA$ , the final cyclization step, formation of the new carbon-carbon bond has been shown to occur from the side of the molecule opposite to that of proton removal (4).

After feeding tryptophan (Trp), mevalonic acid and acetate, all bearing <sup>14</sup>C, it was suggested that dimethylallyltryptophan might be an early precursor (5). This suggestion was in keeping with the biosynthetic route of the clavine alkaloids (6). In fact, the route

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<sup>&</sup>lt;sup>2</sup> Abbreviations used are: Trp, tryptophan;  $\alpha$ - and  $\beta$ -cyclopiazonic acid; cAATrp, cyclo-acetoacetyl-L-tryptophanyl; DMAPP, dimethylallylpyrophosphate; IPP, isopentenyl pyrophosphate; CFE, cell-free extract.

is via the transfer of a  $C_5$  moiety from DMAPP to  $\alpha$ -acetyl- $\gamma$ -( $\beta$ -indolyl)methyltetramic acid (cyclo-acetoacetyl-L-tryptophanyl) (cAATrp) (7). This reaction is mediated by the enzyme dimethylallylpyrophosphate:cycloacetoacetyltryptophanyl dimethylallyltransferase which, throughout this paper, will be referred to as secondary transferase ("S") to distinguish it from dimethylallylpyrophosphatetransferase (EC 2.5.1.1.), primary transferase ("T").

This isoprenylation of Trp or a derivative is not novel; in 1972 Allen (8) reported the transfer of the dimethylallyl group to cycloalanyltryptophanyl in position 2 of the indole nucleus in *Aspergillus amstelodami*, while in 1971 the enzyme responsible for  $4-\gamma\gamma$ -dimethylallyltryptophan itself was isolated from *Claviceps* sp. (6), thus reinforcing considerable evidence of this route in the clavine alkaloids.

## **MATERIALS**

Merck silica gel F<sub>254</sub> tlc plates, precoated 0.25-mm thick, were obtained from Merck, Germany. [1-<sup>14</sup>C]Isopentenylpyrophosphate, L-[G-<sup>3</sup>H]tryptophan, L-[methyl-ene-<sup>14</sup>C]tryptophan, L-[carboxyl-<sup>14</sup>C]tryptophan and [2-<sup>14</sup>C]mevalonic acid were all obtained from Amersham, England.

## **SYNTHESES**

Dimethylallylpyrophosphate (DMAPP) and Isopentenylpyrophosphate (IPP)

These were prepared by the general method described by Cornforth and Popják (9) and purified by chromatography on Sephadex G-25 instead of silica gel, which had the advantage of increased loads and permanency and yielded a purer product. The column was  $80 \times 2.6$  cm, equilibrated and eluted with propanol:concentrated NH<sub>4</sub>OH (65:35). The eluate was monitored using silica TLC plates developed by propanol:concd NH<sub>4</sub>OH:0.001 M EDTA (6:2:2), and the phosphates detected with a molybdate spray (10).  $R_f$ 's of the monophosphates and pyrophosphates of the two alcohols were 0.33–0.44 and 0.18–0.24, respectively. Analysis of the pyrophosphates indicated a purity of 95%.

 $4,5^{-14}C]$  Dimethylallylalcohol,  $[4,5^{-14}C]$  Dimethylallylmonophosphate and  $[4,5^{-14}C]$ -Dimethylallylpyrophosphate

Dimethylallylalcohol was prepared by the method of Plieninger and Immel (11) and phosphorylated according to Cornforth and Popják (9).

# [1-14C]Dimethylallylpyrophosphate

It was possible to produce  $[1^{-14}C]DMAPP$  with relatively high specific activities  $(0.1-60.0 \ \mu\text{Ci}/\mu\text{mole})$  from commercial  $[1^{-14}C]IPP$  by using isopentenyl pyrophosphate isomerase (EC 5.3.3.2) obtained from pig liver (12). The conversion incubation was:  $[1^{-14}C]$ isopentenyl pyrophosphate, 100 nmoles, 6  $\mu$ Ci; Tris-acid maleate, pH 6.0,

50  $\mu$ moles; MnSO<sub>4</sub>·7H<sub>2</sub>O, 10  $\mu$ moles; isomerase, 2.5 mU (1 mU will produce 1 nmole of DMAPP in 1 min at 30°C); and H<sub>2</sub>O to 1.0 ml. The isomerization was carried out at 30°C for 30 min and stopped by immersion in boiling water for 3 min. After cooling, 10  $\mu$ mole of DMAPP was added to the solution and the denatured protein removed by centrifugation. Addition of a few drops of concd NH<sub>4</sub>OH to the supernatant precipitated Mn(OH)<sub>2</sub> after 16 hr. The mixture of isomers could be separated chromatographically (13). The two isomers were detected by scanning for <sup>14</sup>C, CMAPP had an  $R_f$  of 0.35–0.40 and IPP, 0.50–0.55. Integration of the peaks gave a ratio DMAPP: IPP of 2:1 in several experiments. This was confirmed by acidifying a sample (hydrolyzing DMAPP to methylvinylcarbinol) saturating with MgSO<sub>4</sub>·7H<sub>2</sub>O and extracting with toluene several times. This figure is lower than those reported by Holloway and Popják (12) and Shah et al. (13) who quote  $K_{eq}$  of 5 and 6.7, respectively.

The DMAPP was eluted from the paper using 0.1 M NH<sub>4</sub>OH and taken to dryness. Sufficient Tris-acid maleate (pH 8.0, 1 mM) was added to give a DMAPP concentration of 10  $\mu$ moles/ml.

## $\alpha$ -Acetyl- $\gamma$ -( $\beta$ -indolyl)-methyltetramic acid (cyclo-acetoacetyl-L-tryptophanyl)

Five hundred milligrams of [G- $^3$ H]Trp, 408  $\mu$ Ci/mmole was treated with 0.38 ml of thionyl chloride in 5.0 ml of methanol. The solution was kept at 0°C for 16 hr and 20°C for 1 hr before removing the reagents with a stream of dry  $N_2$ . The dried residue, dissolved in 5 ml of methanol, was treated at room temperature for 15 min with sodium methoxide (60 mg Na in 0.8 ml methanol). The mixture was cooled to  $-10^{\circ}$ C, treated with 0.3 ml of diketene and stirred at 20°C for 20 hr, then filtered and concentrated. The concentrate was separated on silica gel to give N-acetoacetyltryptophan methyl ester (604 mg), 550 mg of which was dissolved in 15 ml of benzene, treated with sodium methoxide (40 mg Na/5 ml methanol) and refluxed for 50 min. The mixture was poured on ice and extracted with benzene. On acidification, the aqueous phase was extracted into chloroform to give  $\alpha$ -acetyl- $\gamma$ -( $\beta$ -indolyl)-methyltetramic acid, generally  $^3$ H labeled in Trp (450 mg), mp 167–168°C (from benzene). Anal. Found: C, 63.85; H, 6.20; N, 15.70.  $C_{15}H_{14}N_2O_3$  requires: C, 64.14; H, 6.01; N, 15.57%.

 $\gamma\gamma$ -4-Dimethylallyl-[G- $^3H$ ]tryptophan. We thank Professor C. W. Holzapfel of the Randse Afrikaanse University, Johannesburg, for a generous gift of this compound.

#### METHODS

#### Protein Determination

Protein was determined by the method of McGrath (14), modified to contain half the reported concentration of cyanide.

## Microbiological

P. cyclopium Westling strain 1082 was grown at 25°C in shake culture (200 rpm) in a synthetic medium (4).

## Radioactive Feeding and Determination

The time and duration of feeding depended on the specific activity of the precursor. If this was low, then a pool of the labeled precursor was desirable over a long period of growth. Results then emphasize dilution of specific activity (i.e., specific activity of precursor/specific activity of metabolite). If specific activity was high and the concentration low, then a shorter feeding time, just past the maximum rate of secondary metabolism, was desirable, and the emphasis placed on total incorporation. The precursors were fed as aqueous solutions sterilized by filtration.

Radioactivity was determined using a Packard Tri Carb spectrometer with Triton X-100:toluene (1:2) liquid scintillation mixture (15).

In the case of double-labeling experiments, the isotopes were separated and estimated using a Packard sample oxidizer (Model 306).

Paper and silica chromatograms were scanned using a Berthold scanner (LB 2722).

## Isolation and Estimation of $\alpha$ - and $\beta$ -Cyclopiazonic Acids

These metabolites were isolated from the acidified, homogenized mycelium and filtrate mixture by extraction with chloroform. This was extracted with 0.1 M ammonia which, in turn, was acidified and reextracted into chloroform. The residue was made up to a standard volume containing about 20 mg of the cyclopiazonic acids per ml of chloroform (estimated by uv,  $\log \varepsilon = 4.33$  at 280 nm) and 1 ml applied to a silica gel tlc plate  $(20 \times 20 \text{ cm})$  which was developed using ethyl acetate: methanol: concd NH<sub>4</sub>OH (85:15:10). The metabolites were located by uv ( $\alpha$ CA and  $\beta$ CA,  $R_f = 0.53$  and 0.40, respectively) as very narrow absorbing bands then scraped off and eluted with methanol.

The methanolic solution of  $\alpha$ - or  $\beta$ CA from the above tlc partition was applied to a silica gel tlc plate which had been dipped in an 8% (w/v) methanolic solution of oxalic acid and air dried. The plate was developed using chloroform: methylisobutylketone (4:1) (16). The  $\alpha$ CA ( $R_f$ , 0.7) and  $\beta$ CA ( $R_f$ , 0.5) were detected by uv, scraped off, and eluted with chloroform, to be estimated spectrometrically and radiometrically.

## Cell Breakage and Cell-free Extract (CFE)

The mycelium was washed with buffer (Tris-chloride, pH 7.9, 10 mM; mercapto-ethanol, 2 mM; sodium diethyldithiocarbamate, 2 mM), packed into a press, frozen in Dry Ice and broken by ice shear (17). The broken mycelium was extracted with four volumes of the buffer by stirring for 15 min at  $5^{\circ}$ C. The CFE was obtained by centrifuging at 48 000 g for 1 hr at  $2^{\circ}$ C, followed by careful separation of lipids by decantation and filtration through a large-diameter filter paper.

## Cell-free Biosynthesis of $\beta CA$

A 20-ml reaction mixture containing: L-[G- $^3$ H]Trp, 4  $\mu$ moles, 1  $\mu$ Ci; [1- $^{14}$ C]-DMAPP, 2.5  $\mu$ moles, 0.75  $\mu$ Ci; Tris-chloride, 1 mmole, pH 7.9; CFE, 10 ml, was incubated at 30°C for 1 hr. It was stopped by immersion in boiling water for 3 min and the denatured protein centrifuged off.

To the supernatant was added sufficient Sephadex G-25 (fine) so that an immobile

mass was formed. This was mobilized by the addition of a minimal amount of propanol:  $0.1 M NH_4OH (70:30)$  and loaded onto a Sephadex G-25 (fine) partition column equilibrated with the above propanol solution.

The denatured protein was processed as shown in Scheme 1 (Fig. 1).

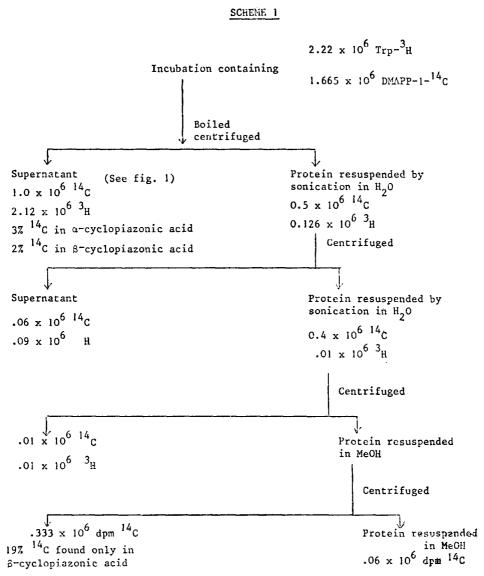


Fig. 1. Treatment of denatured protein from cell-free biosynthesis of  $\beta$ CA. Figures are disintegration per minute for the two isotopes.

## Assay of Secondary Dimethylallyltransferase ("S")

The incubation consisted of: Tris-acid maleate, 50 μmoles, pH 7.2; [1-14C]DMAPP, 100 mmoles, 80 000 dpm; cAATrp, 100 nmoles, and 1-5 mUnits of enzyme in a total

RADIOACTIVE PRECURSOR INCORPORATION INTO A-CYCLOPIAZONIC ACID TABLE 1

Expt. No.*	Compound	Feeding (day-on-day-off)	Total radioactivity (10 <sup>-6</sup> dpm)	Specific activity (mµCi/µmole)	Specific activity of Total Specific cyclopiazonic radioactivity activity acid (10 <sup>-6</sup> dpm) (mμCi/μmole) (mμCi/μmole)	Dilution <sup>6</sup> of specific activity	Yield (#moles per 300 ml)	Incorporation (%)
-	L-tryptophan, methylene-14C-labeled	25	8.44	12.6	1.25	10	357	12
7	DL-tryptophan, methyiene-14C-labeled	2-5	13.10	9.7	2.78	3.5	357	17
33	No addition	Į	1	1	1	}	946	1
4	Dimethylallyl pyrophosphate, 4,5-14C-labeled	25	2.90	4.0	0.17	23	1351	17.6
2	Dimethylallyl monophosphate, 4,5-14C-labeled	2-5	6.90	4.0	0.03	123	893	6.0
9	Dimethylallyl alcohol, 4,5-14C-labeled	2-5	6.45	4.0	0.02	200	952	7.0
7	Mevalonic acid, 2-14C-labeled	2-5	16.80	18.6	0.83	22	946	10.0
∞	y, y-Dimethylallyl tryptophan (br-[G-³H]Trp)	\$	2.02	234.0	0.028	8300	251	0.5
6	γ, γ-Dimethylallyl tryptophan (pt[G-³H]Trp)	4-5	2.02	234.0	No. <sup>3</sup> H	ł	164	1
	Tryptophan, methylene-14C-labeled		22.20	54 500	5.2	10 480		8.4
10	Tryptophan, methylene-14C-labeled	4-5	22.20	2000	5.9	338	208	12.2

" Experiments 1-7 used one inoculum and identical conditions producing 1.3  $\pm$  0.1 g mycelium/flask; three flasks per experiment each containing 100 ml of medium. Experiments 8-10 used one inoculum different from previous group producing the same dry weight of mycelium.

\* Dilution of specific activity = specific activity of precursor/specific activity  $\alpha CA$ .

of 1.0 ml. A blank reaction was always included where the enzyme was replaced by  $H_2O$ . The incubation was for 10 min at 30°C and was stopped by the addition of 2 drops of concd HCl. It was then extracted with benzene, 1.0 ml containing 10  $\mu$ g of  $\beta$ CA. A 100- $\mu$ l aliquot of the benzene layer was applied to a tlc silica plate and developed using ethylacetate: methanol: concd NH<sub>4</sub>OH (85:15:10). The  $\beta$ CA,  $R_f = 0.40$ , was scraped directly into a scintillation vial and counted for 10 min after cooling for at least 2 hr.

### Amino Acid Pools

These were measured on cultures grown from one inoculum. Cultures were grown as described and harvested at 12-hr intervals (10 flasks were grown per time interval). Crude extracts (uncentrifuged) were prepared as described for CFE and aliquots taken to estimate the amino acid pools.

The aliquot was boiled under reflux while stirring with chloroform: methanol (3:1). The solvents were removed by centrifugation, and the residue extracted twice with water and the supernatants mixed. The pH was adjusted to 2.0 with sulfosalicyclic acid and centrifuged. The clear aqueous material was adjusted to pH 7.0 before being taken to dryness. The residue was dissolved in 50 ml of lithium citrate buffer pH 2.2, before loading an aliquot onto a Technicon ion-exchange column (Chromo-beads, Type B)  $0.6 \times 140$  cm. Elution was performed using a gradient in both pH and counter ion (Li<sup>+</sup>) running from pH 3.0 to 6.8. The temperature was kept at 37°C for 5 hr and then at 71.5°C for 16 hr (Technicon Instruction Manual AAA-1).

# RESULTS AND DISCUSSION

## Radioactive Precursor Experiments

The production of  $\alpha$ CA is apparently increased by addition of DMAPP and decreased by Trp (Table 1), a fact often noted during our work. In experiments 1 and 10 it seems that less Trp brings about a greater inhibition; however this anomalous result is due to a variation in the ratios of  $\alpha$ - and  $\beta$ CA which can occur from inoculum to inoculum. Tables 2 and 3 confirm the effect of DMAPP and Trp on the total  $\alpha$ - and  $\beta$ CA production, respectively.

Culture age (days)	CA (µmoles/300 ml)	Mycelial weight (g/300 ml)	CA (µmoles/g mycelium)
2	286 (214)	3.3 (3.2)	84 (66)
3	678 (595)	3.9 (3.6)	174 (161)
4	1250 (893)	4.2 (3.6)	298 (241)
5	1339 (893)	4.3 (3.9)	311 (228)

<sup>\* 1.0</sup> mM added to medium before inoculation.

<sup>&</sup>lt;sup>b</sup> Values in parentheses are controls, i.e., without exogenous DMAPP.

	TABLE 3
THE EFFECT OF TRP	ON THE PRODUCTION OF $\alpha$ - AND $\beta$ CA
	(Total)

Culture age (days)	$\alpha$ - and $\beta$ CA ( $\mu$ moles/300 ml)	Mycelial weight (g/300 ml)
1	36 (36)	0.6 (1.2)
2	214 (223)	2.4 (3.0)
3	381 (804)	2.7 (3.4)
4	402 (1250)	3.1 (3.7)
5	625 (1340)	3.6 (4.4)

 $<sup>^{\</sup>rm a}$  5.0 mM Trp added to the media (3  $\times$  100-ml) before inoculation.

DMAPP is also a better precursor or intermediate than either the parent alcohol or the monophosphate. In fact, what incorporation there is in the latter two cases, probably occurs via other routes because they were fed over a long time. The other suspected intermediate, dimethylallyltryptophan (5), was not incorporated at all when mixed with Trp, though when fed alone it exhibited 0.5% incorporation which was probably due to its catabolism. The dependence of "Dilution of specific activity" on the concentration is well illustrated in experiments 1, 2, 9 and 10. It is obvious in 9 and 10 that Trp is highly diluted by the endogenous system.

Because of the failure of dimethylallyltryptophan to be incorporated and also because Trp inhibited  $\alpha$ CA synthesis, it was felt that a catabolite of Trp could be an intermediate. However, Table 4 indicated that all the C atoms of Trp were incorporated directly into  $\alpha$ CA.

 $\begin{tabular}{ll} TABLE~4\\ Tryptophan~Incorporation~into~\alpha\mbox{-Cyclopiazonic}~Acid \\ \end{tabular}$ 

Compound	Feeding (day-on-day-off)	Radioactivity (µCi)	<sup>3</sup> H/ <sup>14</sup> C in cyclopiazonic acid (dpm × 10 <sup>-5</sup> )	Ratio	
L-[G- <sup>3</sup> H]tryptophan L-tryptophan, methylene-		222.0	11.73/1.2	9.8	
<sup>14</sup> C-labeled L-[G- <sup>3</sup> H]tryptophan	3–4	22.2 222.0	10.44/0.94	11.0	
t-tryptophan, carbonyl- 14C-labeled	3–4	22.2			

#### Amino Acid Pools

Because Trp is such an important precursor, its endogenous pools were measured over the period of production of  $\alpha$ Ca. Table 5 indicates that it exists, if at all, as an

<sup>&</sup>lt;sup>b</sup> Values in parentheses are controls, i.e. without exogenous Trp.

TABLE 5
AMINO ACIDS

Age (hr)	36	48	60	72	84
Dry weight (g) <sup>a</sup>	3.6	6.3	8.7	9.7	11.8
Soluble protein (mg) <sup>a</sup>	424	719	870	408	490
$\alpha$ - and $\beta$ CA (mmoles) <sup>a</sup>	0.15	0.30	0.60	1.31	1.37
Amino acid	Micro	omoles per gra	m dry weight	of mycelium	
Cysteic acid	1.1	1.45	1.45	1.40	2.5
Aspartic acid	5.1	3.3	2.1	1.8	2.0
Threonine	2.9	3.3	2.0	2.3	3.8
Serine	4.0	5.8	3.2	4.7	3.8
Asparagine	0	0	0	0	0
Glutamic acid	20.0	18.2	11.8	7.7	2.7
Glutamine	39.4	24.6	13.6	17.2	13.5
Proline	1.8	2.9	1.9	1.6	2.2
Glycine	3.3	3.6	2.0	2.4	2.2
Alanine	48.8	24.6	7.6	9.8	10.6
Cysteine	0	0	0	0	0
Valine	6.5	10.5	6.3	2.3	5.7
Methionine	1.1	1.8	0.5	0.4	0.7
Isoleucine	1.5	2.9	1.4	1.6	1.8
Leucine	3.6	1.7	4.1	3.9	4.6
Tyrosine	1.5	2.2	1.3	1.4	1.8
Phenylalanine	1.8	3.3	2.1	2.2	2.5
β-Alanine	0	0	0	0	0
y-Aminobutyric acid	25.4	52.2	18.0	22.7	27.0
Tryptophan	0	0	0	0	0
Ornithine	1.5	1.8	0.7	1.2	1.2
Lysine	3.3	4.8	3.7	3.2	2.9
Histidine	1.1	1.5	1.5	1.2	1.6

<sup>&</sup>quot; Per liter of culture.

extremely small pool compared to the other amino acids. The other main feature of this table is the preponderance of glutamic acid, glutamine, alanine, and  $\gamma$ -aminobutyric acid which, together, make up more than 50% of the total amino acids at any time.

## Cell-free Biosynthesis of \( \beta C A \)

The outstanding incorporation of both Trp and DMAPP into  $\alpha$ CA in whole culture made it imperative to test the CFE for similar activity. Figure 2 shows the distribution of the two isotopes in the supernatant. About 10% of the total <sup>14</sup>C fed is shown in Fig. 2; 30% was bound to a protein which precipitated on boiling (Fig. 1) and 55% was eluted as DMAPP around tube 200 which is not shown in Fig. 2.

Tubes 23-45 when taken to dryness lost 90% of the <sup>14</sup>C. Because of its volatility, it was thought to be an alcohol derived from DMAPP. The remaining isotopes of tubes 23-45 were dissolved in methanol and the cyclopiazonates isolated by tlc as described

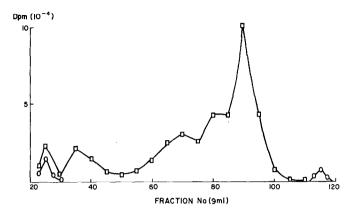


Fig. 2. Partition of supernatant from cell-free biosynthesis of  $\beta$ CA after heat denaturation. Eluant was propanol: 0.1 M NH<sub>4</sub>OH (7:3) on a column (100 × 2.5 cm) of Sephadex G-25 (fine). 9-ml fractions were collected and  $^{14}$ C(0) and  $^{3}$ H( $\square$ ) measured. [1- $^{14}$ C]DMAPP is not shown, eluting around tube 200.

in Methods. No <sup>3</sup>H was found in either  $\alpha$ - or  $\beta$ CA, but the former possessed 0.15% and the latter 0.1% of the original <sup>14</sup>C.

The denatured protein was found to bind 30% of the total <sup>14</sup>C present in the CFE incubation, 19% (still of the total <sup>14</sup>C) of which was shown to be incorporated only into  $\beta$ CA; there was no <sup>3</sup>H found in  $\beta$ CA while  $\alpha$ CA was devoid of any radioactivity (Fig. 1).

The  $\beta$ CA separated from the denatured proteins (Fig. 1) had 5 mg authentic  $\beta$ CA added to it and was isolated using in sequence the tlc systems described (Methods), thus confirming the product as  $\beta$ CA.

It was clear that the CFE contained both an enzyme and a substrate which, along with DMAPP, would produce  $\beta$ CA. This co-substrate was, therefore, a derivative of Trp, which we showed to be  $\alpha$ -acetyl- $\gamma$ -( $\beta$ -indolyl)methyl tetramic acid (7). An incubation similar to that described under Assay of secondary dimethylallyltransferase ("S") using [G-3H]cAATrp as well as [1-14C]DMAPP, yielded  $\beta$ CA with a <sup>3</sup>H: <sup>14</sup>C, ratio which indicated a 1:1 reaction of the substrates (Table 6). This is in keeping with the pathway shown in (Fig. 3) Scheme 2.

TABLE 6

Incubation of [G-3H]cAATrp (100 nmoles, 32 000 dpm) and [1-14C]DMAPP (100 nmoles, 80 000 dpm) with 1 ml of Cell-Free Extract<sup>4</sup>

Product		activity pm)	Substrate in product (nmoles)			
	<sup>3</sup> H	14C	cAATrp	DMAPP		
βCA	781	1845	2.44	2.30		

<sup>&</sup>quot; See enzyme assay in Methods for details of incubation.

The route from Trp to cAATrp is inoperative in the CFE system described, otherwise [G- $^3$ H]Trp would have appeared in  $\beta$ CA. Trp was readily metabolized into two new more nonpolar substances (fractions 70 and 80, Fig. 2), both of which total more than

Fig. 3. The pathway of cyclopiazonic acid biosynthesis.

the remaining Trp. In view of Gatenbeck and Sierankiewiczs' work (19) on tenuazonic acid, another tetramic acid, it is possible that these two metabolites might be N-acetyl-tryptophan and N-acetoacetyltryptophan.

## Variations of cAATrp with Age

cAATrp is present maximally on day 4 and is found predominantly in the culture filtrate (Table 7). However, the CFE represents a 1 to 4 dilution of the mycelium and so the concentration of the metabolite in the filtrate and the mycelium is approximately equal, which suggests a process of diffusion rather than active excretion. It perhaps represents an attempt by the cell to excrete Trp.

TABLE 7
Distribution and Pool Variation of cAATrp with Time

				Age of	culture	•		
	2 days		3 days		4 days		5 days	
Boiled CFE (μl)	200	400	200	400	200	400	200	400
<sup>14</sup> C in β-cyclopiazonic acid (dpm)	300	580	600	1170	1200	1900	1050	1500
Boiled culture filtrate ( $\mu$ l) <sup>14</sup> C in $\beta$ -cyclopiazonic acid	100 100	200 200	100 1500	200 2200	100 2000	200 3000	100 1600	200 2750

Incubation and assay are described in Methods, cAATrp being replaced by boiled CFE or culture filtrate.

## Secondary Dimethylallyltransferase

This enzyme is found only in the mycelium. It shows a direct relationship between enzyme concentration and  $\beta$ CA production (Table 8), over a period of 10 min (Table 9) with an excess of both substrates. The pH profile is broad, with an optimum lying between 6.9 and 7.5 (Table 10).

Protein (μg) (3-day cell-free extract)  14C in β-cyclopiazonic acid	0	20	40	80	160	240
(dpm)	0	440	810	1380	2230	3030

<sup>&</sup>quot; Incubation and assay are described under Methods.

TABLE 9  $\label{eq:table 9}$  Time Dependence of  $\beta$ -Cyclopiazonic Acid Production<sup>4</sup>

Time (min)	0	3	6	10	20	30	40
<sup>14</sup> C in cyclopiazonic acid (dpm)	90	634	1140	1670	2070	2170	2070

<sup>&</sup>lt;sup>a</sup> Incubation is described in Methods.

TABLE 10
pH° Dependence of the Secondary Transferase

рH	5.4	5.9	6.5	6.9	7.5	7.9	8.5
<sup>14</sup> C in β-cyclopiazonic acid (dpm)	450	1810	2220	2680	2570	2480	238

<sup>&</sup>quot;Buffer: Tris-acid maleate at 0.05 I throughout pH range. See Methods for assay.

Presumably this secondary transferase is in direct competition with prenyltransferase (EC 5.3.3.2) for the common substrate DMAPP. The branch point thus formed will control the routing of DMAPP into polyisoprenoids (sterols, ubiquinones etc.) or secondary metabolites, providing that the enzymes are in excess and available to the same pool(s) of substrates.

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