

Studies on the Biosynthesis of γ -Decalactone in *Sporobolomyces odorus*

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Dedicated to Professor Friedrich Drawert on the occasion of his 65th birthday

Incubation of racemic ethyl [2,2- $^2\text{H}_2$]-(*E*)-3,4-epoxydecanoate and the corresponding acid with intact cells of *Sporobolomyces odorus* led to the formation of deuterium labeled γ -decalactone. The detection of labeled 2-decen-4-olide and 4-oxodecanoic acid made it possible to propose the biogenetic sequence leading to γ -decalactone. The determination of the enantiomeric ratio of the lactone revealed an unspecific metabolism of the precursor. In contrast to the (*E*)-isomer, after administration of ethyl [5,6- $^2\text{H}_2$]-(*Z*)-3,4-epoxydecanoate no transformation could be detected. These results indicate that (*E*)-3,4-epoxydecanoic acid, formed from (*E*)-3-decenoyl-CoA, an intermediate of the β -oxidation of linoleic acid, is the genuine precursor in the biosynthesis of γ -decalactone.

Introduction

Although aliphatic γ - and δ -lactones are important compounds, *e.g.* as flavor constituents in foodstuffs and as attractants in pheromone research [1–3], their biogenetic formation still awaits elucidation. The identification of presumable precursor compounds as trace constituents in the spectrum of plant volatiles [4] and the detection of various enzymatic activities by incubation experiments with fruit tissue made it possible to propose some biogenetic pathways [5, 6].

Glycerol bound 4- and 5-keto acids, identified as trace constituents in milk fat [7, 8], have been discussed as potential precursor compounds of γ - and δ -lactones in fermented milk products. The reduction of 4- and 5-keto acids by microorganisms are well known transformations [9], but the formation of these keto acyl glycerols in milk fat remained unclear.

Walker and Dimick postulated a δ -oxidation of saturated fatty acyl CoA-esters after acetate derived *de novo* synthesis of the corresponding acids [10, 11]. The first intermediate of this pathway, 4-enoyl CoA, might be enzymatically hydrated to the 4- or 5-hydroxy derivative, the immediate biogenetic precursor of the lactones.

When [1- ^{14}C]decanoic acid was administered to cells of the lactone producing red yeast *S. odorus*, γ -decalactone contained 84% of the total radioac-

tivity in the extracts of volatiles [12]. But, corresponding to the results of Dimick and Walker, no intermediates were detected, which made it possible to confirm the mechanistic pathway.

The increase in concentration of γ -dodecalactone and (*Z*)-6-dodecen-4-olide in the milk of ruminants after application of a diet with a high degree of oleic and linoleic acid [13] indicated that these lactones might derive from the degradation of long chain fatty acids.

Detailed physiological studies on *S. odorus* revealed that the degradation of previously biosynthesized fatty acids corresponds with an enhanced accumulation of γ -nona-, γ -decalactone and (*Z*)-6-dodecen-4-olide [14].

(*Z,Z*)-3,6-dodecadienoyl-CoA, (*Z*)-4-decenoyl-CoA and (*E,Z*)-2,4-decadienoyl-CoA are intermediates of the β -oxidation of linoleic acid [15, 16]. However, no hydration of these possible precursors leading to the corresponding 4-hydroxyacyl intermediates could be demonstrated in previous investigations [17, 18].

Based on these results and on the fact that cytochrome P-450 dependent monooxygenases catalyze the hydroxylation of different compounds, *e.g.* fatty acids, hydrocarbons and steroids [19], we assumed that a comparable enzymatic system is responsible for the formation of aliphatic lactones. In incubation experiments with cells of *S. odorus* using various deuterated fatty acids, such as (*Z*)-4-decenoate, (*E*)-3-decenoate and decanoate, no transformation neither into γ - nor into δ -decalactone could be detected [20].

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For the purpose to trace the metabolic routes leading to γ -decalactone the capacity of submerged strawberry tissue to transform racemic [2,2- $^2\text{H}_2$]-(*E*)-3,4-epoxydecanoic acid into the lactone could be demonstrated [14]. The optical purity of the product, however, did not correspond to the naturally formed optical pure (*R*)-(+)- γ -decalactone.

In this paper the biogenetic pathway of γ -decalactone is proposed based on the results obtained after GC-MS identification of metabolites and chiral analysis of the lactone formed after incubation of deuterium labeled (*E*)-3,4-epoxydecanoic acid.

Materials and Methods

Chemicals

(*E*)-3-decenoic acid was obtained from Ferak (Berlin, F.R.G.), (*E*)-2-heptenal, ethyl acrylate, and triphenylphosphine were purchased from Aldrich (Steinheim, F.R.G.), D_2O and NaOD (30% in D_2O) were from Roth (Karlsruhe, F.R.G.); pig liver esterase (EC 3.1.1.1.) 260 U/mg protein from Sigma (Deisenhofen, F.R.G.); (*R*)-(+)-1-phenylethylisocyanate (PEIC), *N,N*-4-dimethylaminopyridine and *m*-chloroperbenzoic acid (mCPBA) from Fluka (Neu-Ulm, F.R.G.) and 2-decenoic acid was a gift from Haarmann & Reimer (Holzminden, F.R.G.).

Syntheses

Ethyl [2,2- $^2\text{H}_2$]-(*E*)-3,4-epoxydecanoate

1 g (*E*)-3-decenoic acid, 200 μl NaOD and 3 ml D_2O were heated in a gastight reaction vessel for 24 h at 170 °C. After acidification and extraction, the product was esterified with ethanol/acetylchloride (9:1).

EI-MS (Ethyl [2,2- $^2\text{H}_2$]-(*E*)-3-decenoate): *m/z* 200 (M^+ , rel. int. 0.8%), 56 (100), 43 (84), 41 (76), 57 (69), 55 (67), 69 (67), 110 (63), 90 (60), 70 (58), 71 (53); Kovats-Index (IK) DB Wax 60 W: 1680.

950 mg mCPBA (= 5.5 mmol), dissolved in 2.5 ml diethylether were added to 0.93 g ester (= 5 mmol), dissolved in 5 ml ether at 0 °C. After 48 h, the solvent was evaporated and the product was cleaned up by silica gel liquid solid chromatography (LSC) (glass column, 80 \times 1.5 cm). The

elution was performed with 150 ml pentane/ CH_2Cl_2 (2:1), 150 ml pentane/ether (9:1) and 150 ml pentane/ether (5:1). 20 ml fractions were collected and analyzed by gas chromatography.

EI-MS (Ethyl [2,2- $^2\text{H}_2$]-(*E*)-3,4-epoxydecanoate): *m/z* 216 (M^+ , rel. int. 0.02%), 55 (100), 41 (62), 103 (48), 43 (46), 56 (42), 69 (33), 57 (32), 73 (31), 75 (27), 45 (25); IK (DB WAX 60 W): 1958.

^1H NMR (270 MHz, CDCl_3): 2.76 ppm (1 H, dt, $J_{\text{H}_3-\text{H}_4} = 2.4$ Hz; $J_{\text{H}_4-\text{H}_5} = 5.4$ Hz); 3.03 ppm (1 H, m (br.)).

[2,2- $^2\text{H}_2$]-(*E*)-3,4-epoxydecanoic acid

The ester bound was hydrolyzed with commercially available pig liver esterase. 100 mg ethyl [2,2- $^2\text{H}_2$]-3,4-epoxydecanoate were dispersed in 15 ml phosphate buffer pH 7.0. After addition of 50 μl esterase solution (= 0.5 mg protein) the reaction mixture was shaken for 6 h at 25 °C leading to a quantitative hydrolysis; the yielded acid was extracted with 100 ml diethylether.

Ethyl (*E*)-3,4-epoxydecanoate

(*E*)-3-decenoate was esterified and epoxidized as described.

EI-MS (Ethyl (*E*)-3-decenoate): *m/z* 198 (M^+ , rel. int. 2%), 55 (100), 41 (95), 69 (82), 43 (63), 54 (48), 110 (45), 88 (42), 83 (33), 39 (32).

EI-MS (Ethyl (*E*)-3,4-epoxydecanoate): *m/z* 185 ((*M*-29), rel. int. 0.1%), 55 (100), 41 (98), 43 (97), 101 (75), 71 (58), 57 (54), 56 (52), 42 (45), 69 (44), 73 (39).

^1H NMR (270 MHz, CDCl_3): 2.54 ppm (2 H, dd, $J_{\text{H}_2-\text{H}_2'} = 16.2$ Hz; $J_{\text{H}_2-\text{H}_3} = 5.7$ Hz); 2.75 ppm (1 H, dt, $J_{\text{H}_4-\text{H}_3} = 2.4$ Hz; $J_{\text{H}_4-\text{H}_5} = 5.4$ Hz); 3.04 ppm (1 H, dt, $J_{\text{H}_3-\text{H}_2} = 5.7$ Hz; $J_{\text{H}_3-\text{H}_4} = 2.4$ Hz).

Ethyl [5,6- $^2\text{H}_2$]-(*Z*)-3,4-epoxydecanoate

1 ml (*E*)-2-heptenal was hydrogenated with Pd, 5% on charcoal and D_2 in a gastight reaction vessel at 10 bar. Purification by fractionated distillation and by LSC on silica gel yielded in 720 μl [2,3- $^2\text{H}_2$]-heptanal.

EI-MS ([2,3- $^2\text{H}_2$]-heptanal): *m/z* 116 (M^+ , rel. int. 0.5%), 45 (100), 43 (55), 42 (54), 44 (53), 71 (52), 72 (42), 46 (42), 41 (41), 56 (32), 83 (10); IK (DB WAX 60 W): 1135.

According to the method described [21], 1.626 g triphenylphosphine and the aldehyde were combined and heated at 125 °C. 650 μ l ethyl acrylate were added to this clear solution and refluxed for 24 h. The reaction product, ethyl [5,6-²H₂]-3-decenoate (yield 30%, ratio (*Z*):(*E*) = 88:12), was purified by silica gel LSC. The resolution of the (*Z*)/(*E*) isomers was achieved by LSC on silica gel, impregnated with Ag(NO₃) [22].

EI-MS (Ethyl [5,6-²H₂]-(*Z*)-3-decenoate): *m/z* 200 (M⁺, rel. int. 0.2%), 55 (100), 41 (99), 70 (86), 56 (74), 42 (72), 88 (57), 111 (51), 153 (17), 154 (8); IK (DB WAX 60 W): 1641.

Epoxidation and purification of the epoxide were performed as described for the (*E*)-isomer.

EI-MS (Ethyl [5,6-²H₂]-(*Z*)-3,4-epoxydecanoate): *m/z* 171 (M-45, rel. int. 0.2%), 56 (100), 55 (93), 43 (84), 101 (73), 71 (73), 42 (72), 41 (67), 57 (63), 44 (47), 70 (45); IK (DB WAX 60 W): 1980.

¹H NMR (270 MHz, CDCl₃): 2.55 ppm (2H, dd, *J*_{H2-H2'} = 14 Hz; *J*_{H2-H3,H2'-H3} = 6.3 Hz); 2.97 ppm (1H, dt, *J*_{H4-H3} = 2.5 Hz, *J*_{H4-H5} = 6.3 Hz); 3.30 ppm (1H, dt, *J*_{H3-H4} = 2.5 Hz, *J*_{H3-H2} = 6.3 Hz).

Feeding experiments

S. odorus ATCC 24259 was grown aerobically (air flow rate: 1 vvm) at 24 °C in a 10 l – fermenter. The medium contained (in 1 l of deionized water): 45 g sucrose, 5 g lactose, 3.0 g MgSO₄ · 7 H₂O, 2.5 g (NH₄)₂SO₄, 2.5 g KH₂PO₄, 2.5 g L-Alanin and 0.1 g CaCl₂ · 2 H₂O. After 77 h 1 l culture broth (= 3 g cell dry weight) was centrifuged for each incubation experiment and the cells were washed twice with m/15 phosphate buffer, pH 6.8. Cells were resuspended in 30 ml phosphate buffer and filled in 100 ml Erlenmeyer flasks, containing 10 mg of the incubation substrate. The flasks were shaken at 26 °C for 4 h and the reaction was stopped by adding 50 ml diethylether with subsequent extraction of volatiles.

Analysis of volatiles

The incubation broth was extracted with 2 × 50 ml diethylether and the acids were methylated with diazomethane. After concentration to 0.2 ml, the extract was transformed to a LSC column (4.5 g silica gel) and separated into four 40 ml-fractions: (Fraction I: pentane/CH₂Cl₂

(9:1); fraction II: pentane/CH₂Cl₂ (2:1); fraction III: pentane/ether (9:1); fraction IV: ether).

Gas chromatographic analysis was performed on a Carlo Erba 4100 gas chromatograph, equipped with a DB Wax 60 W (60 m/0.32 mm internal diameter, J & W).

Determination of optical purity

γ -Decalactone was isolated by preparative gas chromatography. As described elsewhere [23], the lactone was transformed with butylamine into the corresponding 4-hydroxydecanamide. The conversion into the diastereomeric 4-[(*R*)-[(1-phenylethyl)carbamoyl]oxy]N-butyldecanamide was achieved by reaction with (*R*)-(+)-1-phenylethylisocyanate.

Mass spectrometry

GC-MS investigations were carried out using a Finnigan (quadrupole) mass spectrometer, model 4510, combined to a Finnigan 9610 gas chromatograph, equipped with a DB Wax 60 W. The diastereomeric 4-[(*R*)-[(1-phenylethyl)carbamoyl]oxy]N-butyldecanamides were separated on a DB 210 capillary column (30 m/0.32 mm internal diameter, J & W).

For electron impact ionization (EI-MS) the ionization potential was 70 eV and the scan range was 35–350 Da/0.8 s. For chemical ionization (CI-MS) methane was used as the CI reagent gas at a pressure of 0.5 torr. The source temperature was maintained at 200 °C. The scan range was 80–350 Da/0.8 sec. Scanning and data processing were accomplished with an INCOS data system.

¹H NMR Spectroscopy

¹H NMR spectra were recorded at 270 MHz on a Bruker WH 270 NMR spectrometer in CDCl₃ solutions. Chemical shifts are referenced to tetramethylsilane ((CH₃)₄Si) as internal standard.

Results

Incubation substrates

Fig. 1 presents the spectral data of ethyl [2,2-²H₂]-(*E*)-3,4-epoxydecanoate and the corresponding non labeled compound. The characteristic fragmentation pattern of the CI spectra makes

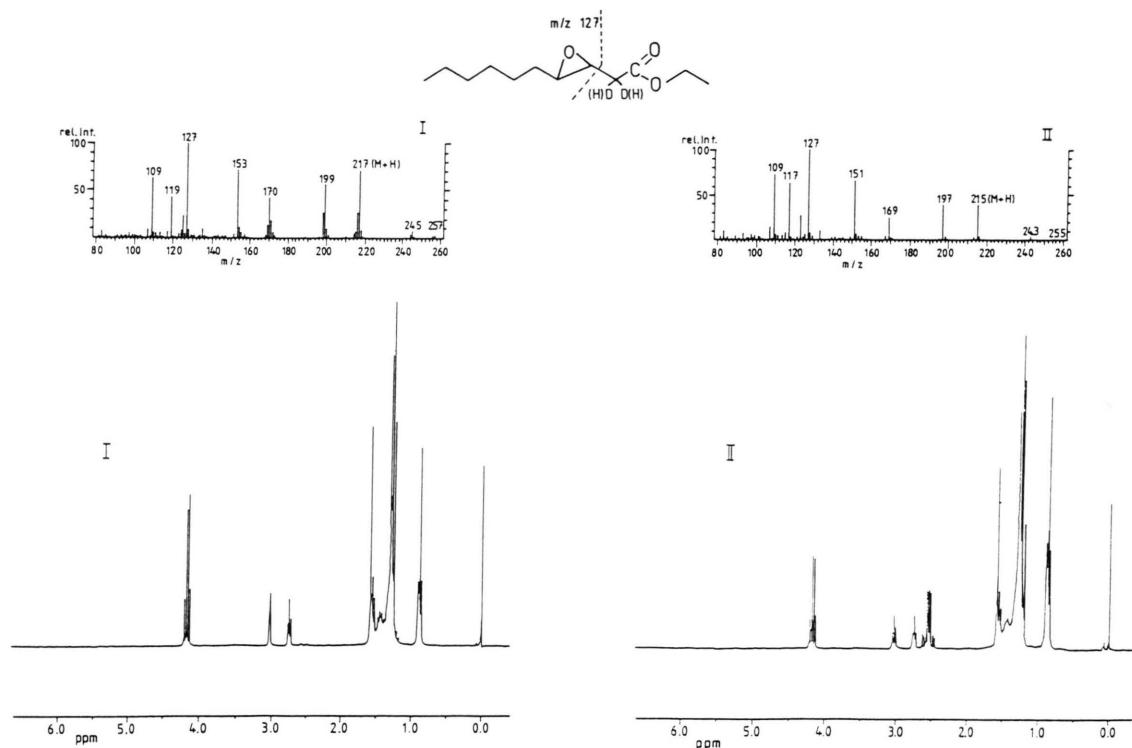


Fig. 1. CI-MS and ¹H NMR spectra of ethyl [2,2-²H₂]-(*E*)-3,4-epoxydecanoate (I) and ethyl (*E*)-3,4-epoxydecanoate (II).

it possible to localize the position of the isotopes. The base peak (m/z 127) in both compounds refers to the cleavage between C₂ and C₃. In conclusion, the deuterium atoms are fixed to C₂. The ¹H NMR spectra are in accordance with the mass spectra. Compared to the unlabeled reference, the resonance signal for the α -methylene protons disappeared and the double triplet for the proton at C₃ shifted to a broad multiplett, due to the coupling of ¹H-²H.

The chemical stability of the substrates was tested under incubation conditions. After extraction and chromatography on silica gel neither a hydrolytic cleavage of the oxirane ring nor a loss of deuterium could be observed.

The mass spectrum of ethyl [5,6-²H₂]-(*Z*)-3,4-epoxydecanoate reveals that the substrate was not uniformly deuterated in position C₅ and C₆ (Fig. 2). Both, the [M + H]⁺-peak (m/z 216/217) and the fragment ions m/z 128 and 129, resulting from the α -cleavage indicate that the compound

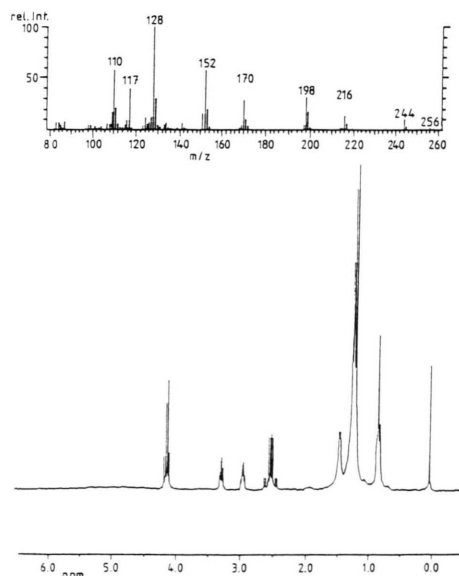


Fig. 2. CI-MS and ¹H NMR spectra of ethyl [5,6-²H₂]-(*Z*)-3,4-epoxydecanoate.

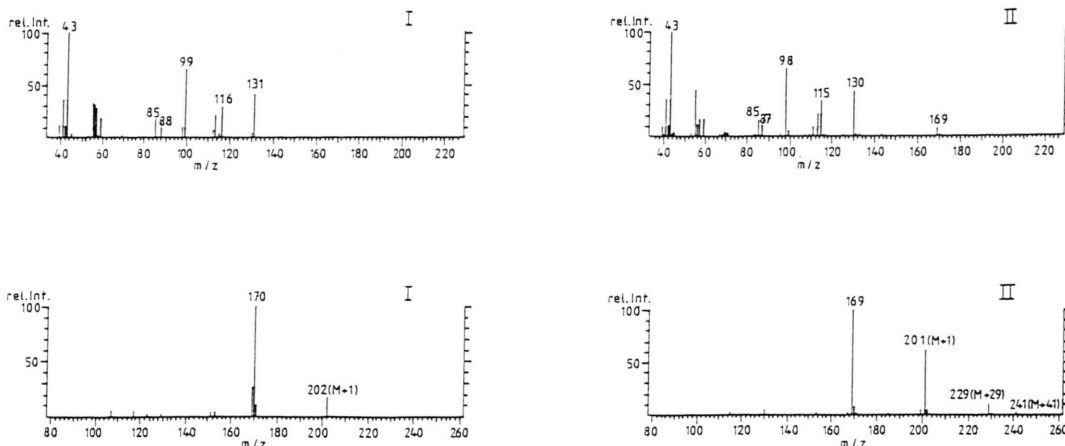


Fig. 3. EI- und CI-MS of methyl 4-oxodecanoate: (I) detected in fraction III of the ether extract after incubation of ethyl [2,2- $^2\text{H}_2$]-(*E*)-3,4-epoxydecanoate and (II) authentic reference.

was a mixture of single and double deuterated epoxy ester.

Feeding experiments

Different incubation experiments revealed that added precursors were optimally metabolized by cells harvested after a fermentation period of 60 to 80 h. Additionally, the low concentrations of naturally formed fatty acids and lactones makes it possible to detect the labeled products even at low metabolization rates of the added incubation substrates.

Incubation of ethyl (*E*)-3,4-epoxydecanoate and (*E*)-3,4-epoxydecanoic acid

A high esterase activity of the *S. odorus* cells resulted in a complete hydrolysis of the added ethyl ester within a incubation period of 4 h. The metabolization rate of the hydrolytically formed epoxy acid was about 5%, comparable to the result obtained after incubation of (*E*)-3,4-epoxydecanoic acid.

The metabolites subsequently described were detected after incubation either of the ester or of the acid.

The ether extract of the incubation broth was methylated and fractionated on silica gel before GC and GC/MS investigation. The non-metabolized substrate and methyl 4-oxodecanoate were collected in Fraction III. The deuterium labeling of methyl 4-oxodecanoate is evident by its mass

spectral data (Fig. 3). In the EI mass spectrum, the fragment ions m/z 116 (α -cleavage), m/z 131 (McLafferty rearrangement) and m/z 99 (m/z 131 – CH_3OH) contain the α -methylene group and suggest the presence of only one deuterium. Additionally, molecular weight determination by CI-MS ($[\text{M} + \text{H}]^+$: m/z 202) points to the loss of one ^2H -isotope in the course of generation of this compound. An enzymatic step should be responsible for the abstraction of one deuterium, because the chemical stability of the labeling has been verified.

The reconstructed ion chromatogram of the polar fraction IV of the ether extract is presented in Fig. 4. Peak a, b, and c correspond to the naturally

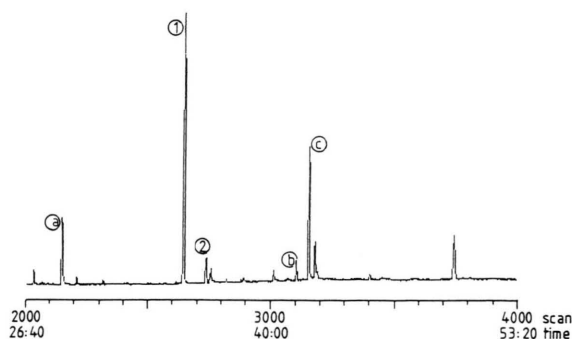


Fig. 4. Capillary gas chromatogram (reconstructed ion detection) of the polar fraction after incubation of ethyl [2,2- $^2\text{H}_2$]-(*E*)-3,4-epoxydecanoate to cells of *S. odorus* (DB WAX 60 W 60 m \times 0.32 mm i.d., temperature programmed from 70 $^\circ\text{C}$ (3 min isothermal) to 220 $^\circ\text{C}$ at 4 $^\circ\text{C}/\text{min}$).

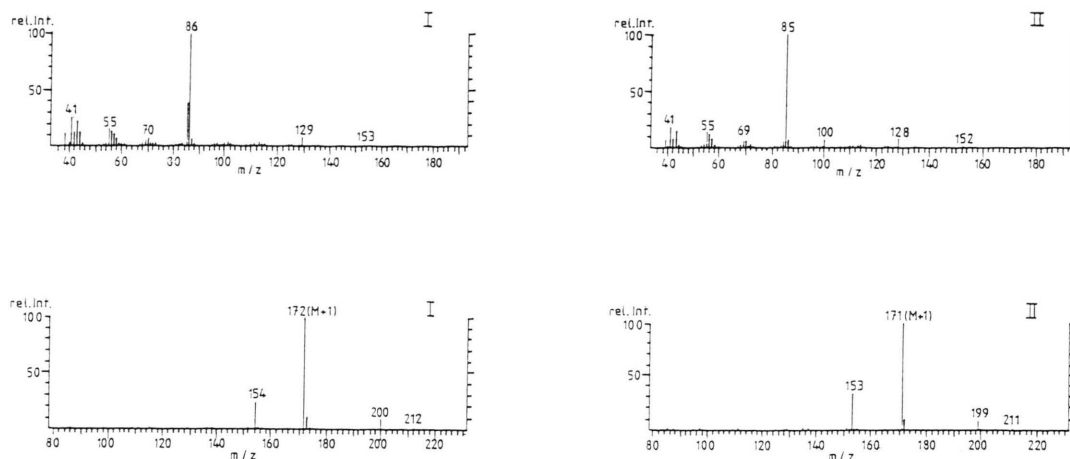


Fig. 5. EI- and CI-MS of γ -decalactone: (I) corresponding to peak I in Fig. 4; (II) authentic reference.

formed, unlabeled metabolites 2-phenylethanol (a), γ -dodecalactone (b), and (*Z*)-6-dodecen-4-olide (c), respectively.

Fig. 5 shows the mass spectrum of γ -decalactone (Peak I in Fig. 4). The base peak at m/z 86 instead of m/z 85 and the molecular weight determination ($[M + H]^+$; m/z 172 instead of m/z = 171) characterize the difference of the labeled and unlabeled compound.

The enantiomeric composition of the formed γ -decalactone, isolated by preparative GC, was investigated by capillary GC after derivatization into the corresponding diastereomeric 4-[(*R*)-(1-phenylethyl)carbamoyloxy]N-butyldecanamides [23]. The chromatograms of the separated diastereoisomers, presented in Fig. 6, correspond to the racemic reference (I), the naturally biosynthesized lactone (II), and to the product, isolated from the incubation experiment (III). The enantiomeric ratio of the lactone, isolated after the biotransformation of the epoxide, was (*S*):(*R*) = (31:69)%, compared to the biosynthesized optically pure (*R*)-(+)- γ -decalactone; this indicates a non-enantioselective metabolism of the added racemic precursor. Additional experiments revealed that the enantiomeric ratio is not constant but depends on (a) the transformation rate of the added precursor and (b) on the amount of previously biosynthesized lactone in the course of growth.

The mass spectral investigation of the separated diastereoisomers gives evidence for the assumption of an unspecific metabolism (Fig. 7). The

mass spectra of 4-[(*R*)-(1-phenylethyl)carbamoyloxy]N-butyldecanamides are characterized by the fragment ion $M-164$, according to the abstraction of phenylethylcarbamate. Therefore, the fragment m/z 227 of the derivative of the (*S*)-enantiomer reveals the deuterium labeling. The derivative of the

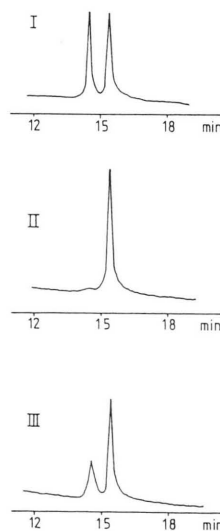


Fig. 6. Capillary GC-separations of (*S*)- and (*R*)-4-[(*R*)-(1-phenylethyl)carbamoyloxy]N-butyldecanamide (DB 210 30 m \times 0.32 mm i.d.; temperature programmed from 220 $^{\circ}$ C (5 min isothermal) to 260 $^{\circ}$ C at 1 $^{\circ}$ C/min): (I) derivatives of racemic γ -decalactone; (II) derivatives of biosynthesized lactone of *Sp. odorus*; (III) derivatives of γ -decalactone, isolated after incubation of ethyl [2,2- 2 H $_2$]-(*E*)-3,4-epoxydecanoate.

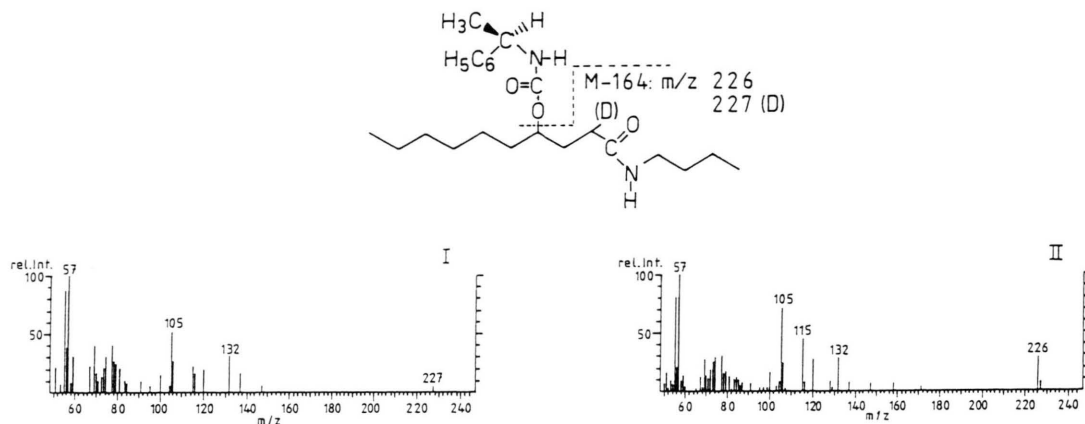


Fig. 7. EI-MS of the separated diastereomeric derivatives of γ -decalactone, isolated from the polar fraction of the extract after incubation of ethyl (*E*)-3,4-epoxydecanoate: (I) corresponding to (*S*)- γ -decalactone; (II) corresponding to (*R*)- γ -decalactone.

(*R*)-enantiomer, with m/z 226 and m/z 227 is a mixture of the naturally formed and the precursor derived lactone.

Peak 2 in the reconstructed ion chromatogram corresponds to 2-decen-4-olide, which was identified by comparison to gas chromatographic and mass spectrometric data of the synthesized unlabeled reference compound (Fig. 8). The fragment ion m/z 85 in the labeled metabolite, in contrast to m/z 84 of the authentic reference, is due to the abstraction of the alkyl moiety after hydrogen rearrangement [24]. After administration of unlabeled ethyl (*E*)-3,4-epoxydecanoate, the mass

spectra of 2-decen-4-olide obtained from the incubation experiment were identical to the reference.

*Incubation of ethyl [5,6- $^2\text{H}_2$]-(*Z*)-3,4-epoxydecanoate*

In contrast to the described biotransformation of the (*E*)-isomer, after incubation of ethyl [5,6- $^2\text{H}_2$]-(*Z*)-3-decenoate none of the described metabolites were detected. The EI- and CI mass spectra of γ -decalactone, which has been the main constituent of the biotransformation of the (*E*)-isomer, correspond exactly to the reference com-

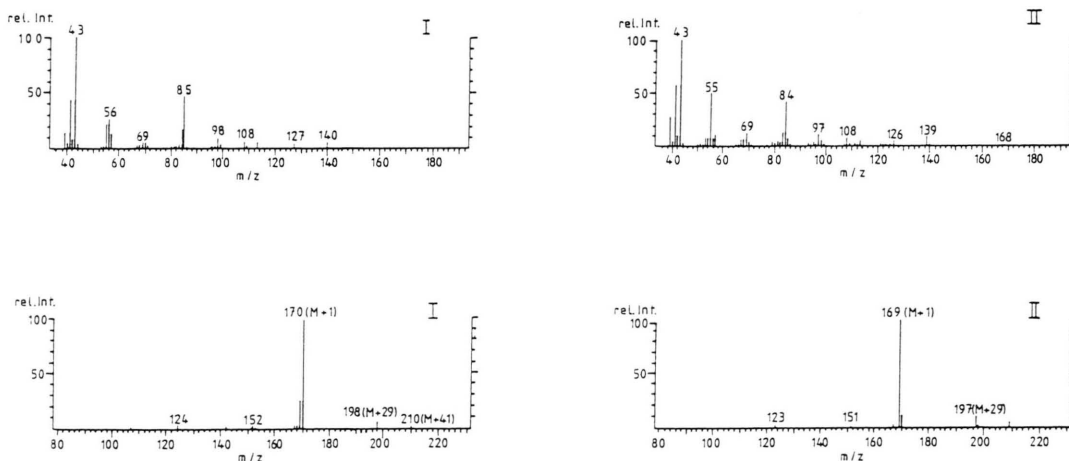


Fig. 8. EI- and CI-MS of 2-decen-4-olide: (I) corresponding to peak 2 in Fig. 4; (II) authentic reference.

pound. Apart from the hydrolysis of the ethyl ester, no additional reaction could be detected.

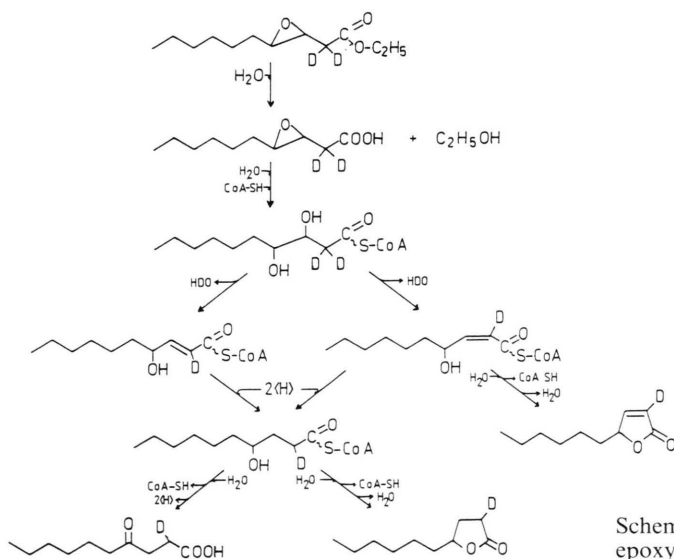
Discussion

Scheme 1 summarizes the proposed enzymatic steps leading to the identified products after incubation of ethyl [2,2- $^2\text{H}_2$]-(*E*)-3,4-epoxydecanoate or the corresponding acid. Generally, epoxides are metabolized by hydration, catalyzed by epoxide hydrolases (EC 3.3.2.3.) or glutathione S-transferases (EC 2.5.1.18.). In plant and microbial systems, only the hydrolases have been characterized. According to their physiological function as detoxifying enzymes, hydrolases from microsomes of animal tissue catalyze the hydration of a broad spectrum of substrates [25, 26]. In contrast to that, but corresponding to the specificity of the hydrolysis of the epoxy group observed in this study, the substrate specificity of an epoxide hydrolase, responsible for the biosynthesis of trihydroxystearate in the skin of young apples is very high [27].

The first intermediate in the metabolism of the epoxide, 3,4-dihydroxydecanoic acid, could not be detected. We assume that after intracellular hydration, the polar dihydroxyacid can not pass the membranes. Alternatively, it might be possible that the intermediate dihydroxyacid is rapidly oxidized with high enzymatic activity. Studies on the

formation of lactones from long chain fatty acids using biotransformations of hydroxylated substrates, e.g. ricinoleic acid and erythro-9,10-dihydroxystearic acid, resulted also in an oxidation to the corresponding oxo or ketohydroxy compounds, respectively [20]. Analogously, the postulated intermediate might be oxidized to 3-oxo-4-hydroxydecanoic acid, which is chemically unstable.

The accumulation of 2-decen-4-olide can be interpreted by two enzymatic steps with subsequent chemical or enzymatically catalyzed lactonization. After transformation of the dihydroxyacid into its CoA derivative, the formation of 4-hydroxy-2-decenoyl-CoA can be explained by a reverse reaction of an enoyl-CoA hydratase. Recently, Smeland *et al.* described the activities of two enoyl hydratases with opposite enantioselectivities, responsible for the epimerization of (*R*)-3-hydroxyoctanoyl-CoA [28]. An unspecific hydration of the racemic epoxy ester is leading to a dihydroxy acid with both (*R*)- and (*S*)-3-hydroxy groups. According to the described mechanism [28], the (*R*)-3-hydroxy group can be dehydrated to 4-hydroxy-(*E*)-2-decenoyl-CoA or into 4-hydroxy-(*Z*)-2-decenoyl-CoA. Because the formation of 2-decen-4-olide requires the (*Z*)-configuration of the double bond, the reverse reaction of the (*Z*)-2-enoyl hydratase should be active. The detection of this unsaturated lactone, which is not



Scheme 1. Proposed transformation of ethyl (*E*)-3,4-epoxydecanoate by cells of *S. odorus*.

naturally formed by *S. odorus* indicate that the epoxidation of (*E*)-3-decenoic acid or its CoA-ester proceeds enantioselectively. Additionally, this reaction explains the loss of one deuterium in all labeled metabolites.

After enzymatic hydrogenation, the resulting 4-hydroxydecanoic acid, which could be detected as a trace constituent, either cyclizes to the final metabolite, γ -decalactone, or is oxidized to 4-oxodecanoic acid.

The formation of 2-alken-4-olides and the corresponding saturated lactones with chain length from C₆ to C₁₀ by the basidiomycete *Polyporus durus* [29] can be also explained with the proposed pathway.

After incubation of (*Z*)-3,4-epoxydecanoic acid, no metabolites could be detected. Therefore, we assume that the (*E*)-isomer is the genuine precursor leading to γ -decalactone. (*E*)-3-decenoyl-CoA is formed during β -oxidation of linoleic acid after hydrogenation of (*E,Z*)-2,4-decadienoyl-CoA [17, 18]. The first step in the biosynthesis of γ -decalactone in *S. odorus* should be the epoxidation of the double bond of this intermediate. The fact, however, that this initial step could not be demonstrated after administration of deuterated (*E*)-3-decenoic acid, can be explained by the sensitivity of the enzymes involved in the epoxidation. Duppel *et al.* demonstrated that yeast microsomal

hydroxylation systems are sensitive towards higher concentrations of substrates [30]. Because of the high optical purity of the naturally occurring γ -decalactone and the non-enantioselective transformation of (*E*)-3,4-epoxydecanoate, it can be concluded that the biogenesis is initiated by an enantioselective epoxidation.

In contrast to γ -decalactone, other γ -lactones which are accumulating in the fermentation broths of this yeast, γ -hexa-, γ -nona-, and (*Z*)-6-dodecen-4-olide are not optically pure but occur in constant enantiomeric ratios [20]. The minor enantioselectivity during formation of these compounds may be referred to the activity of other enzymes, which exhibit different selectivities, or a common enzymatic system, responsible for the formation of all lactones, characterized by a substrate dependent enantioselectivity.

For a more detailed investigation of this enzymatic system, it is necessary to reproduce the described metabolic steps in a cell-free extract of this organism. But, until now, the disintegration of the yeast cells resulted in a complete loss of the capacity to metabolize added epoxides.

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