



Review: Biocatalytic transformations of ferulic acid: an abundant aromatic natural product

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In this review we examine the fascinating array of microbial and enzymatic transformations of ferulic acid. Ferulic acid is an extremely abundant, preformed phenolic aromatic chemical found widely in nature. Ferulic acid is viewed as a commodity scale, renewable chemical feedstock for biocatalytic conversion to other useful aromatic chemicals. Most attention is focused on bioconversions of ferulic acid itself. Topics covered include cinnamoyl side-chain cleavage; nonoxidative decarboxylation; mechanistic details of styrene formation; purification and characterization of ferulic acid decarboxylase; conversion of ferulic acid to vanillin; O-demethylation; and reduction reactions. Biotransformations of vinylguaicol are discussed, and selected biotransformations of vanillic acid including oxidative and nonoxidative decarboxylation are surveyed. Finally, enzymatic oxidative dimerization and polymerization reactions are reviewed.

Keywords: ferulic acid; biocatalysis; microbial and enzymatic biotransformations

Introduction

Biotechnology has been defined as 'The application of scientific and engineering principles to the processing of materials by biological agents to provide goods and services' [19]. Biocatalysis is a more focused term defining the use of enzymes or of living catalysts such as whole microbial cells for the production of useful chemicals. As in chemical processing, the biocatalyst initiates or accelerates a reaction between two or more substances while the catalyst itself remains unchanged and is recycled. Biocatalysis encompasses the field of microbial transformations of organic or inorganic compounds which result in a change in chemical structure. Useful biocatalytic reactions yield products (often called metabolites) which accumulate in the incubation medium. Biocatalytic processes are now used for the preparation of very valuable specialty chemicals for the pharmaceutical, agricultural, flavor and fragrance, nutritional and chemical industries. Such processes are also exploited for the preparation of commodity chemical substances on an enormous industrial scale.

In this review, we attempt to describe biocatalytic technologies and how they can be used with an extremely abundant, naturally occurring phenolic compound known as ferulic acid. We review the history, isolation and physical properties of ferulic acid and related phenolic cinnamic acids. The widespread occurrences and functions of ferulic acid in nature are described along with several emerging fields concerned with the antioxidant and putative chemopreventative properties of this family of compounds. The development of needed biocatalytic approaches toward the production of useful aromatic/phenolic chemicals from

ferulic acid is also emphasized. The aim of producing valuable chemical products from ferulic acid is coupled with a description of what is known about biotransformation and biodegradation pathways of ferulic acid. Finally, we emphasize detailed efforts from our own laboratories devoted to unraveling mechanistic details associated with the biocatalysis of ferulic acid and its products.

Ferulic acid

Ferulic acid (**1**, in Figure 1), 3-(4-hydroxy-3-methoxyphenyl)-2-propenoic acid is an extremely abundant and nearly ubiquitous phenolic cinnamic acid derivative. It was described in 1866 as a component of *Ferula foetida* Reg. (*Umbelliferae*) and various salts of the compound were prepared [129]. The name ferulic acid obviously derives from the botanical name of the plant from which it was obtained. In nature, ferulic acid is thought to arise biosynthetically in plants via cinnamic acid [54], which finds its origin in the shikimic acid pathway [35]. Since its discovery, numerous reports on the occurrence, chemistry, synthesis, spectral analysis, possible uses and biological properties have appeared in the literature. The structures of ferulic acid and related compounds have been confirmed by their synthesis, by ¹³C-NMR spectral analysis [77,87] and by X-ray analysis [98]. Chemical and biological properties of ferulic acid were recently summarized in an exhaustive review by Graf [47].

The structure of ferulic acid (**1**) is shown in Figure 1 along with the structures of caffeic acid (**3a**) and isoferulic acid (**3b**) which also are abundant in nature. The unsaturated side-chain of ferulic acid can exist naturally in either *trans* or *cis* conformational structures such as **1** or **2**, where hydrogens at the adjacent 7- and 8-positions are either on opposite or the same sides of the double bond [2]. The double bond is subject to *trans-cis* isomerization [41]. Chlorogenic acid (**4**) is also shown in Figure 1 to illustrate

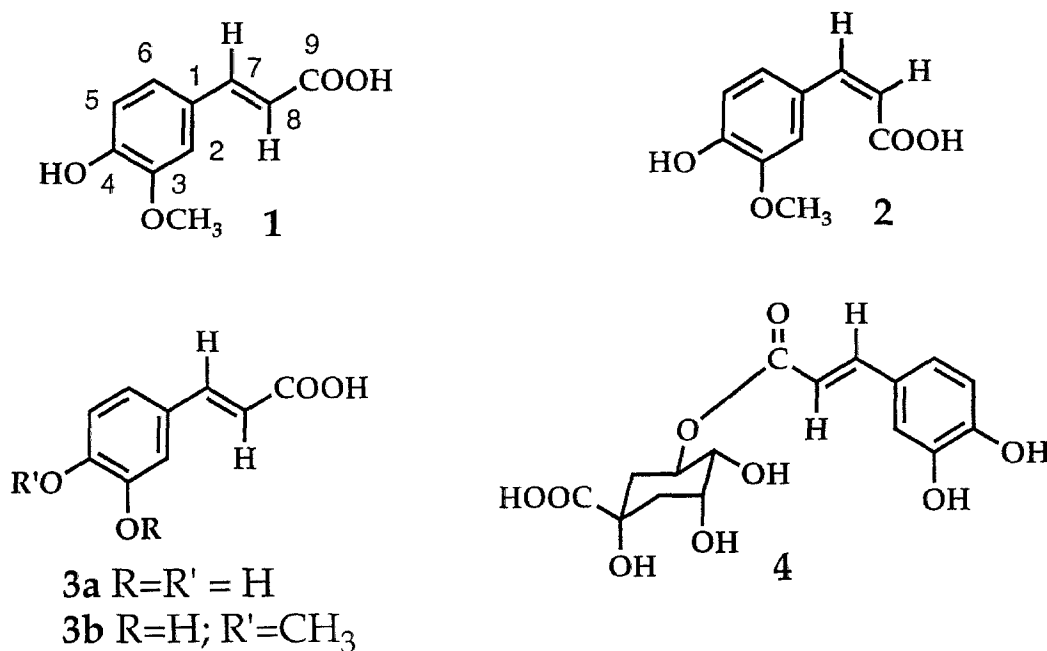


Figure 1 The structures of *trans*-ferulic acid (1), *cis*-ferulic acid (2), caffeic and isoferulic acids (3a, 3b) and chlorogenic acid (4)

the fact that cinnamates may occur free in nature, they may be esterified as with 4, or esterified to hydroxyl groups of polymeric backbones found within the plants that contain them.

In plants, ferulic acid is rarely found in free form [118]. It occurs widely linked to various carbohydrates as glycosidic conjugates, and it occurs as various esters and amides with a wide variety of natural products. As esters, ferulic acid is linked to polysaccharides where it may occur in monomeric or dimeric form [3,46,47]. It also occurs esterified with flavonoids and with various hydroxycarboxylic acids including malic, tartaric, tartronic, hydroxycitric, quinic, gluconic and hydroxy fatty acids [64,65]. Relatively nonpolar ferulate esters occur with long chain alcohols, and with various plant sterols [47]. Ferulic acid is also found esterified with a tetrahydroisoquinoline-mono-terpene glucoside [72], a cyanogenetic glycoside [1,127] and an amino-hydroxycyclopentenone [36]. Levels of ferulic acid amides with putrescine, tyramine or tryptamine may be indicators of environmental stress in plants [90]. Amides of ferulic acid with amino acids or dipeptides are used as preservatives in baking [47]. Corn contains feruloyl tryptamide [37], and ferulic acid esters linked to a polysaccharide [102] and to dihydroxy β -sitosterol [106].

Ferulic acid occurs commonly in fruits, vegetables, grains, beans, leaves, seeds, nuts, grasses, flowers, and other types of vegetation [47,64]. It is abundant in some plants, varying from 1 g kg⁻¹ in corn kernels [102,132] to 0.05 g kg⁻¹ in refined wheat and 0.5 g kg⁻¹ in whole wheat flour [47] to 1.5 g kg⁻¹ in the spice turmeric [64]. Ferulic acid can be isolated in nearly 2% yield (wt/wt) from the hulls of corn kernels obtained from wet milling [10].

Ferulic acid and its derivatives are well recognized as members of a class of phenolic natural antioxidants [47], and many natural and presumed functions and potential uses of these compounds can be attributed to this property.

The roles of ferulic acid in the plants that contain it and in human nutrition and well-being continue to be elaborated [47]. In plants, alleopathic regulation and protection mechanisms are related to ferulic acid content. Ferulic acid endows structural rigidity and strengthens cell wall architecture by cross-linking pentosan chains, arabinoxylans and hemicelluloses, rendering these components less susceptible to hydrolytic enzymes during germination [47,122]. Considerable evidence suggests that ferulic acid is involved in regulating plant growth. In roots, ferulic acid appears to inhibit growth of competing plants. Ferulic acid exhibits a detrimental effect on seedling growth, plant dry weight and leaf expansion [18,125]. High levels of ferulic acid appear to inhibit root growth and the development of root system architecture [16]. High levels of ferulic acid appear to be inversely related to mineral and water uptake [88], water utilization [18], hyphal elongation [130], and hydrolytic enzyme activities of germinating seeds [34]. Ferulic acid and its derivatives confer protection to plants against avian [52], insect [30,86,119], viral [90] and fungal [107] invasion. Increased levels of ferulylamides are found in plants acclimated to herbicide treatment and nutritional depletion [90]. The UV-absorbing properties of ferulic acid confer additional protection to plants.

Microbial transformations of ferulic acid

Microbial and/or enzymatic transformations of ferulic acid and related phenolic cinnamates have been examined from several interesting perspectives [111]. Ferulic acid and its microbial metabolites appear to play roles in plant germination and plant growth [126]. These compounds influence the makeup of microfloral consortia in soil by inhibiting the growth of certain microorganisms while stimulating others [116]. Styrene metabolites from hydroxycinnamates like ferulic acid confer pest-resistance in some plants [61]. Hydroxystyrenes and other aromatic microbial metabolites

of ferulic acid are also important flavor constituents in beer, wine, soy sauce, and fruit juices [12,26,117]. The rumen metabolism of ferulic acid in cows [101], and the possible involvement of antioxidants like ferulic acid and its metabolites in mammalian toxicology are of growing interest [114]. Ferulic acid also has been used as a 'lignin phenolic substrate to model microbial degradations of aromatic compounds from chemical and petroleum industrial wastes [51,67,78]. The great abundance of ferulic acid in nature has also prompted several investigations into the potential for using ferulic acid as a starting material for microbial or enzymatic synthesis of useful aromatic chemical compounds [33,69]. Conversions of ferulic acid to commercially valuable aromatic aldehydes like vanillin have been described [84,85].

Numerous studies on aerobic or anaerobic biotransformations of ferulic acid have been reported [5–8,25,38,42,48,53,55,63,71,73,75,76,91,92,94,96,97,99,101,103,108,115,117,120,123,124,126]. Major pathways of biotransformation by bacteria, fungi and yeasts are summarized in Figure 2, and in Table 1. The major pathways of metabolism can be summarized as: a) reductive, forming saturated side chains, aldehydes and alcohols such as **6**, **12** and **13**; b) cleavage of acetate from the unsaturated side-chain flowing through vanillin (**5**), a major branch point for the formation of other oxidative and reductive products;

c) oxidation of vanillin to vanillic acid (**7**), a central metabolic product yielding guaiacol (**9**) by decarboxylation, protocatechuic acid (**10**) by *O*-dealkylation, and methoxyhydroquinone (**8**) by oxidative decarboxylation; d) other *O*-dealkylation reactions giving caffeic acid (**3a**) directly from ferulic acid; and e) as perhaps the oldest recorded reaction, nonoxidative decarboxylation to form styrenes such as vinylguaiacol (**14**). Further oxidative degradation of the aromatic ring of ferulic acid or its metabolites usually occurs through protocatechuic acid (**10**). Features of the major types of ferulic acid biotransformation reactions are summarized below.

1. Elimination of two carbons from the cinnamoyl side-chain: Elimination of an acetate moiety from the unsaturated cinnamate side-chain is one of the most common pathways of ferulic acid metabolism in bacteria, yeasts and fungi. Different interpretations of the mechanism by which side-chain cleavage gives rise to biodegradation intermediates have been postulated, but relatively limited work has been done with the enzymes catalyzing this transformation reaction. A variety of pseudomonads catalyze the oxidation of **1** to vanillic (**7**) and protocatechuic (**10**) acids [113,124].

Toms and Woods [124] reported that *P. acidovorans* converted both *cis*- and *trans*-ferulic acid to vanillin (**5**) and

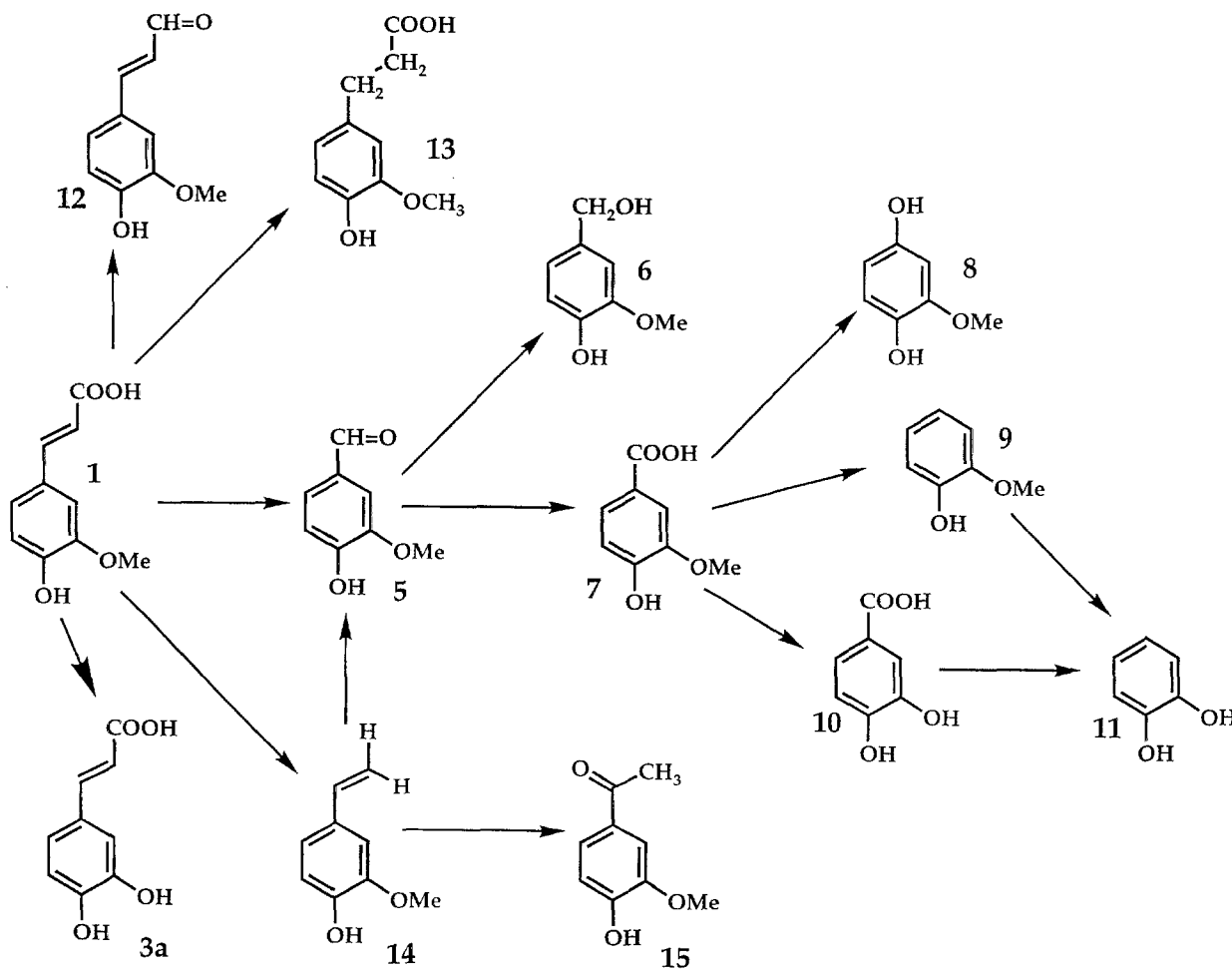


Figure 2 Major pathways of microbial transformation of ferulic acid

**Table 1** A summary of ferulic acid metabolites formed by bacteria, fungi and yeasts

Organism	Compound no, shown in Figure 2:														Ring fission	Ref
	3a	5	6	7	8	9	10	11	12	13	14					
<i>Aerobacter</i> sp															+	[42]
<i>Bacillus megaterium</i>															+	[68,69]
<i>B. pumilus</i> (3) ^a															+	[12,33]
<i>B. subtilis</i>		+		+						+					+	[56]
<i>B. subtilis</i>															+	[68,69]
<i>Corynebacterium glutamicum</i>		+		+								+	+			[84,85]
<i>Enterobacter</i> DG-6	+									+					+	[48]
<i>Erwinia uredovora</i>														+	(caffeic acid)	[61]
<i>Escherichia coli</i>		+		+						+						[103]
<i>Klebsiella oxytoca</i>														+	(caffeic acid)	[61]
<i>Margarinomyces mutabilis</i>					+					+ ^b					+ ^b	[63]
<i>M. heteromorph</i>					+					+ ^b					+ ^b	[63]
<i>Myobacterium</i> sp															+	[66,69]
<i>Nocardia</i> sp															+	[68,69]
<i>Pseudomonas acidovorans</i>		+		+						+						[124]
<i>P. cepacia</i>													+	+		[8]
<i>P. mira</i>		+ ^b		+												[75]
<i>P. putida</i>		+														[84]
<i>P. putida</i> Type A															+	[68,69]
<i>P. sp</i> TMY1009					+					+					+	[113]
<i>Streptomyces rimosus</i>															+	[68,69]
<i>S. setonii</i>		+		+						+						[120]
<i>Wolinella succinogenes</i>					+									+		[101]
Anaerobes from marsh sediment	+										+					[9]
Anaerobic microbial consortium														+		[96]
Methanogenic consortium	+														+	[51]
Rat gastrointestinal microflora														+	+	[114]
Unidentified bacterium					+											[73]
<i>Aspergillus niger</i>															+	[66,68,69]
<i>A. niger</i>		+														[84]
<i>A. carneus</i>															+	[12]
<i>A. ochraceus</i> (2) ^a															+	[12]
<i>A. terreus</i> (2) ^a															+	[12]
<i>Candida intermedia</i>															+	[12]
<i>Corynespora cassiicola</i>															+	[12]
<i>Curvularia affinis</i>															+	[12]
<i>C. clavata</i>															+	[12]
<i>C. lunata</i> (5) ^a															+	[12,68,69]
<i>Fomes fomentarius</i>		+ ^b	+ ^b												+ ^b	[71]
<i>Fusarium coeruleum</i>															+	[12]
<i>F. dimerum</i>															+	[12]
<i>F. eumartii</i>															+	[12]
<i>F. moniliforme</i> (2) ^a															+	[12]
<i>F. oxysporum</i> (5) ^a															+	[12]
<i>F. roseum</i>															+	[12]
<i>F. solani</i> (Mart) Sacc.		+		+						+					+	[97]
<i>F. solani</i> (4) ^a															+	[12]
<i>F. tritinctum</i>															+	[12]
<i>Hansenula anomala</i> (2) ^a															+	[12,68,69]
<i>H. beckii</i>															+	[12]
<i>H. capsulata</i>															+	[12]
<i>H. henricii</i>															+	[12]
<i>H. minuta</i>															+	[12]
<i>H. saturnus</i>															+	[12]
<i>Paecilomyces variotii</i>		+	+	+						+					+	[108]
<i>Penicillium frequentans</i>															+	[68,69]
<i>P. glaucans</i>															+	[66]
<i>P. rubrum</i>	+									+					+	[123]
<i>Pestalotia palmarum</i>															+	[108]
<i>Polyporus versicolor</i>		+ ^b	+ ^b												+	[71]
<i>Pullularia pullulans</i>			+ ^b							+ ^b					+	[63]
<i>Rhizopus arrhizus</i>															+	[68,69]
<i>Rhodotorula rubra</i>					+					+					+	[68,69]
<i>R. sp</i>															+	[126]
<i>Saccharomyces cerevisiae</i> (5) ^a														+	+	[12,68,69,89]
<i>Sporotrichum pulverulentum</i>															+	[20]
<i>Trametes</i> sp			+	+												[99]

^aNumber of strains reported^bProposed only^cFrom vanillic acid

subsequently to vanillic acid (7). The proposed pathway for acetate cleavage from ferulate in the side-chain shortening reaction involves water addition to the double bond to give **16**, subsequent oxidation of **16** to ketone **17** and removal of an acetate unit to give vanillin (**5**) (Figure 3).

Vanillin is further oxidized in a NAD⁺-dependent reaction to form vanillic acid (7). In this work, vanillin and vanillic acid were established as metabolites by infrared spectroscopic comparisons of isolated metabolites to standard compounds. Evidence for the elimination of [2-¹⁴C]-acetate from [2-¹⁴C]-*trans*-ferulic acid supported the proposed mechanism. However, attempts to synthesize the β-hydroxy intermediate (**16**) failed. Quinoid isomerization of the *p*-hydroxyl group on the aromatic ring causes elimination of the benzylic hydroxyl group of **16**. This work demonstrated that microorganisms are clearly capable of oxidizing *trans*-ferulic acid into smaller aromatic metabolites such as vanillic acid (7) common to basic bacterial metabolism.

Although detailed cell-free metabolism experiments were not generally conducted for most organisms listed in Table 1, it is clear that side-chain elimination of an acetate moiety occurs very commonly among bacterial species including *Bacillus* [56], *Corynebacterium* [84,85], *Escherichia* [103], *Pseudomonas* [75,124], *Streptomyces* [120] and other unidentified bacteria [73]; and among fungi and yeasts including species of *Aspergillus* [84], *Fomes* [71], *Fusarium* [97], *Polyporus* [71] and *Rhodotorula* [68,69].

The precise mechanism involved in the cleavage of two carbons from the ferulic acid cinnamoyl side-chain has not been fully established. Intermediates proposed during the cleavage reaction are unstable. Thus, it has not been possible to isolate or characterize them by classical chemical or

biochemical means. With *Rhodotorula rubra*, we employed a combination of biochemical, mass spectral and NMR spectroscopic analytical approaches to probe the side-chain cleavage mechanism [69]. We first established that *R. rubra* catalyzed the conversion of **1** to **7** and later **9** by isolating these metabolites and characterizing them by mass spectrometry (MS) and nuclear magnetic resonance (NMR) analyses. Cell-free extracts were used to demonstrate that the side-chain cleavage reaction was CoA/ATP- and NAD⁺-dependent. Similar results were previously reported with extracts of a *Polyporus* strain, and with *Streptomyces setonii* in ferulic acid transformations to vanillic acid [69,120]. These cofactor requirements suggested the involvement of a classical β-oxidation process similar to fatty acid oxidation, where steps include the hydration of **1** to **16**, oxidation to a ketone (**17**), and subsequent acetate cleavage (Figure 3).

We attempted to confirm this path by incubating *R. rubra* in buffers prepared with 20% H₂¹⁸O [69]. Based on available evidence, intermediates **18–21** (Figure 3) would all be labeled with ¹⁸O. Three different ¹⁸O-labeled species of vanillic acid **22**, **23**, **24** (Figure 3) and unlabeled **7** were obtained from this incubation and analyzed by MS and ¹³C-NMR to measure both the extent of incorporation of labeled oxygen, and the precise location of oxygen in **7**. By MS, unlabeled vanillic acid (**7**, Figure 3) exhibits a molecular ion [M⁺] of *m/z* 168. The intensity of the *m/z* 170 [M⁺ + 2] ion (36%) indicated an equal amount of species of **23** and **24** each of which contained 18% of ¹⁸O. The intensity of the ion at *m/z* 172 [M⁺ + 4] is due to doubly labeled vanillic acid (**22**) containing 3.5% of ¹⁸O. Since **21** contained approximately 20% ¹⁸O, hydrolysis in 20% H₂¹⁸O buffer would theoretically afford doubly labeled **22** in 4% yield.

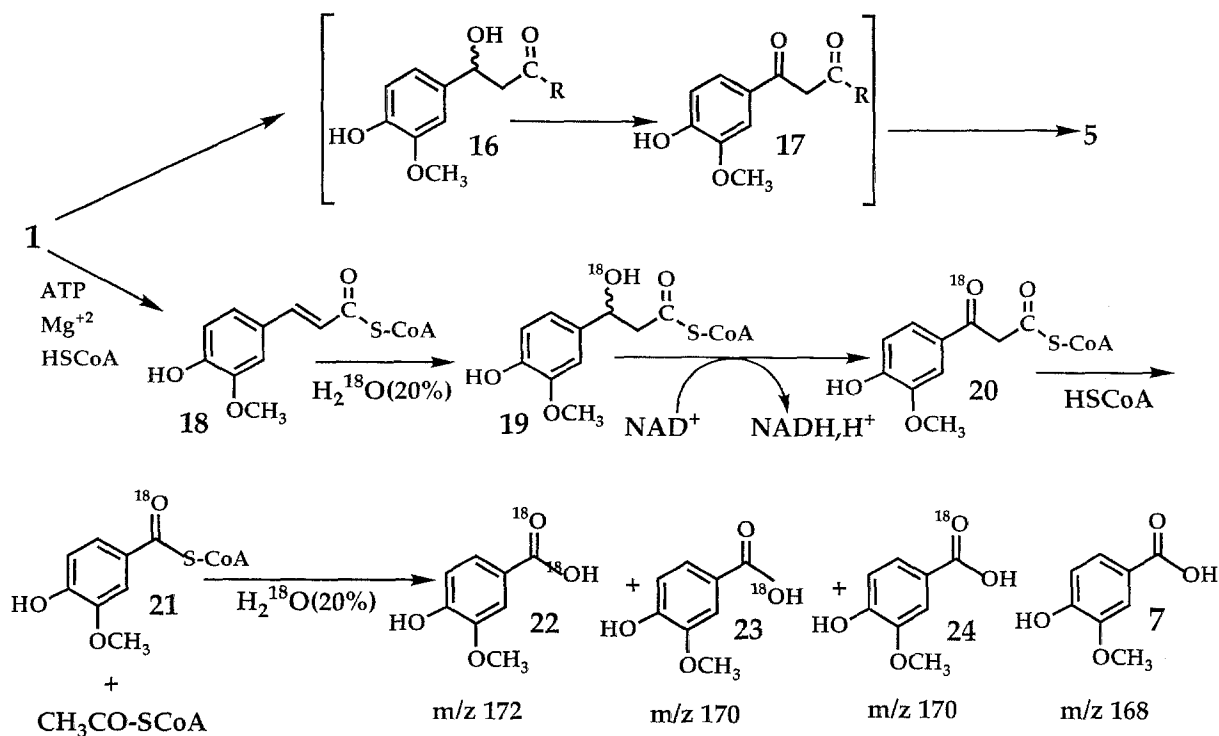


Figure 3 Details of the mechanism by which ferulic acid (**1**) is metabolized to labeled vanillic acids (**22**, **23**, and **24**) and unlabeled vanillic acid (**7**)

The incorporation of ^{18}O specifically into the carboxyl group was confirmed by ^{13}C -NMR where the upfield shift of 2.1 Hz (0.023 ppm) for the carboxyl carbon is caused by the presence of ^{18}O on the carboxyl-carbon atom [69].

2. Nonoxidative decarboxylation of ferulic acid to 4-vinylguaiacol (4-hydroxy-3-methoxystyrene) (14**):** Styrene formation is among the oldest reported biotransformations of cinnamic acid derivatives. In 1906, Herzog and Ripke [66] reported that species of *Penicillium glaucum* and *Aspergillus niger* converted cinnamic acid into styrene. Styrene is also produced from cinnamic acid by other molds [74] and yeast [27,93]. This is interesting because the decarboxylation of cinnamic acid apparently does not require an oxygenated aromatic ring, a common requirement for this biotransformation reaction in most microorganisms [12,42,68,69]. Table 1 reveals that styrene formation is perhaps the most common microbial transformation reaction observed with ferulic acid. The reaction generally occurs rapidly, and in high yield. Although not established with all microorganisms, the decarboxylation reaction appears to be nonoxidative in nature. Scheline observed that rat intestinal microflora convert ferulic acid into 4-vinylguaiacol, thus indicating the potential for this reaction to convert a common dietary chemical into a potentially toxic styrene intermediate in the gastrointestinal tract [114].

Bioconversion of ferulic acid to **14** is often accompanied by additional metabolites. For example, Nazareth and Mavinurke reported the nonoxidative decarboxylation of *trans* ferulic acid to 4-vinylguaiacol by *Fusarium solani* (Mart.) Sacc. 4-Vinylguaiacol was thought to be further metabolized to vanillin, vanillic acid and protocatechuic acid, which undergoes *ortho*-ring cleavage to β -keto adipic acid [97]. Rahouti *et al* also suggested that *Paecilomyces* gave vinylguaiacol as a major metabolite that was converted further to vanillin [108]. However, the amounts of vanillin and vanillic acid observed were very small (probably less than 1% as determined by HPLC), thus casting doubt on this pathway. Samejima *et al* [113] found that *Pseudomonas* TMY1009 produced 4-vinylguaiacol from *trans*-ferulic acid via a proposed 7-hydroxy intermediate such as **16** (Figure 3, R=OH). Decarboxylation of caffeic acid (**3a**, Figure 1) to afford 3,4-dihydroxyphenylethane was thought to involve cinnamic acid double bond reduction and subsequent decarboxylation [134]. This reaction more likely involves initial formation of styrene first and subsequent reduction as described later for biotransformations of 4-vinylguaiacol.

3. Mechanism of styrene production of microorganisms: Surprisingly little work had been conducted until very recently on the mechanism of styrene formation from ferulic acid. Manitto *et al* reported on the stereochemistry of the decarboxylation reaction of 3,4-dimethoxycinnamic acid by cultures of *Saccharomyces cerevisiae* [89]. A deuterium atom on position 8 of the cinnamic acid (^2H , Figure 4) was retained (by ^1H -NMR) in the resulting styrene product. These results were used to suggest that *in vivo* decarboxylation might occur by enzymatic *cis*-addition to the double bond followed by *trans*-decarboxylative elimination

to maintain the observed stereochemistry of the retained deuterium atom.

Our screening studies [68] revealed several microorganisms capable of catalyzing the decarboxylation reaction in good yield. Among these, baker's yeast gave **14** and another metabolite identified as 4-hydroxy-3-methoxyphenylpropionic acid **13** (Figure 2). The side-chain reduction reaction did not occur under aerobic conditions. Reduction of the cinnamoyl side-chain has been observed with bacteria [8,96,101] and fungi [40,68]. An organism originally identified as *Pseudomonas fluorescens* UI-670 also converted **1** to **14** in excellent yield both under standard aerobic incubation conditions, and when the organism was incubated under argon. This organism has since been identified as a strain of *Bacillus pumilus* by fatty acid analysis and metabolic studies (unpublished results). We used this microorganism and *Saccharomyces cerevisiae* to probe the mechanism of decarboxylation further.

A nonoxidative mechanism was envisioned to involve initial enzymatic isomerization of **1** to quinoid intermediate **25** (Figure 4) [68]. If formed, the doubly vinylogous β -keto acid **25** would spontaneously decarboxylate to afford styrene. The mechanism would satisfy the requirement for a *para*-hydroxyl group in the decarboxylation process. Isomerization of **1** to **25** involves addition of a proton from water to the side-chain. Thus, we reasoned that evidence for the existence of **25** could be obtained by conducting the incubation reaction in deuterium oxide-containing media, and by isolating and characterizing the deuterium-labeled styrene **26** by mass spectrometry, and both deuterium and ^1H -NMR. The proposed mechanism is analogous in part to that proposed by Hopper and coworkers and Whited and Gibson for the hydroxylation of cresol by the enzyme cresol methylhydroxylase [133].

Conversions of **1** to **26** were obtained with both baker's yeast and *B. pumilus* UI-670 in media containing 0, 50, 75 or 100% D_2O . Incorporation of deuterium into all samples of **26** isolated from deuterium-containing medium was supported by the presence of a peak at 5.13 ppm in their ^2H -NMR spectra. The absence of any other deuterium peaks by ^2H -NMR indicated that deuterium incorporation was highly stereospecific. The deuterium incorporations were also confirmed by MS where the relative intensities of ions at m/z 150 (unlabeled vinylguaiacol **14**) and m/z 151 (**26**, Figure 4) were compared. All MS analyses indicated that **26** contained a single deuterium atom, the relative intensities of which were proportional to the amounts of D_2O used in incubation buffers. NMR and mass spectral results obtained for the decarboxylation of **1** by baker's yeast were identical to those obtained for *B. pumilus* [68]. These results provided confirmatory evidence for a novel microbial mechanism for the nonoxidative decarboxylation of naturally occurring 4-hydroxycinnamates.

4. Purification and characterization of ferulic acid decarboxylase: Ferulic acid decarboxylase was purified 2780-fold from cells of *B. pumilus* UI-670 by a combination of DEAE-cellulose, hydrophobic interaction chromatography, gel filtration, hydroxylapatite and mono-Q chromatographic steps [70]. The colorless enzyme is constitutive, stable in the cold and does not require exogen-

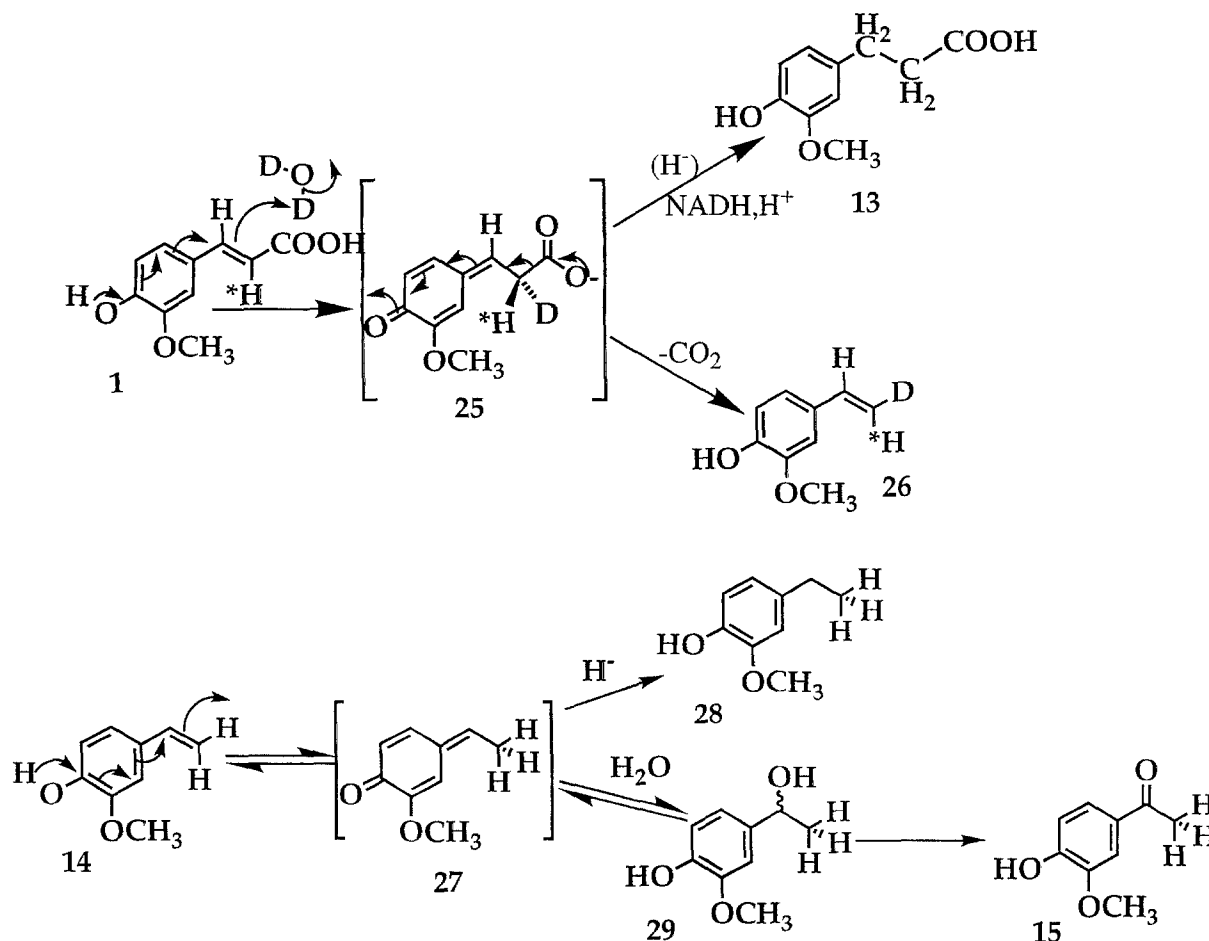


Figure 4 Mechanism of vinylguaiacol formation from ferulic acid in microorganisms, and further oxidation products of vinylguaiacol

ously added cofactors or metal ions for full activity. It exhibits optima at pH 7.3 and 27° C in 20 mM phosphate buffer, and has a K_m of 7.9 mM for ferulic acid. Gel filtration indicated an apparent molecular mass of 40.4 ($\pm 6\%$) kDa whereas Na dodecylsulfate-polyacrylamide gel electrophoresis showed a molecular mass of 20.4 kDa. Thus, it appears that ferulate decarboxylase is a homodimer in solution. The enzyme catalyzed decarboxylations of *p*-coumaric acid, but not 2- or 3-hydroxycinnamic acids, thus indicating the requirement for a 4-hydroxyl group in the decarboxylation reaction. Ferulate decarboxylase contains no apparent flavin component. Further, the reaction was not enhanced by additions of FMN, FAD, NAD⁺, NADP⁺, NADH H⁺, or NADPH H⁺, and it was not coupled to typical electron acceptors. It was inhibited by mercuric and cupric ion, 4-chloromercuribenzoic acid, *N*-ethylmaleimide and iodoacetamide suggesting that sulfhydryl groups are required for catalysis. Lack of inhibition by diethyl pyrocarbonate suggests that histidine is not involved in the decarboxylation reaction.

The N-terminal sequence of 39 amino acids was determined to be:

NH₂-Met-Asp-Gln-Phe-Val-Gly-Leu-His-Met-Leu-Tyr-

15 Thr-Tyr-Glu-Asn-Lys-Trp-Glu-tyr-Glu-Ile-Tyr-Ile-

25 Lys-Asn-Asp-His-Thr-Ile-Asp-Tyr-Arg-Ile-His-Ser-

39 Gly-Met-Val-Gly-

Some of the properties of our purified enzyme were similar to those described earlier for a partially purified *p*-coumarate decarboxylase from *Cladosporium phlei* [58], and, interestingly, for a vanillate hydroxylase (decarboxylating) enzyme from *Sporotrichum pulverulentum* and a vanillyl-alcohol oxidase from *Penicillium simplicissimum* [23]. *p*-Cresol-methylhydroxylase, which is mechanistically similar to our ferulate decarboxylase, is classically a ferrohemo-chromagen oxidoreductase [133]. This purified yellow enzyme contains a covalently bound flavin with absorption maxima at 454 and 344 nm in alkaline solution [133].

In January 1995, Degrossi *et al* reported the purification of ferulate and *p*-coumarate decarboxylase from *B. pumilus* [33]. The enzyme was induced in this culture, was slightly higher in molecular mass (23 kDa by Na dodecyl sulfate PAGE, 45 kDa by size exclusion), showed different pH (pH 5.5), buffer and temperature (37° C) optima and

kinetic properties (K_m 1.03 mM for ferulic acid). A 17-amino acid N-terminal sequence was identical to our ferulate decarboxylase for amino acids 1–9, but differed for amino acids 10 (Ile), 14 (Gln) and 16 (Gly). Interestingly, the decarboxylase reaction is inhibited when our enzyme is incubated in Tris buffer, the buffer used in the first isolation steps of purification of ferulate and *p*-coumarate decarboxylase from *B. pumilus* [33].

Ferulate decarboxylases can be useful biocatalysts for the production of styrenes that in themselves have value as starting materials for the preparation of aromatic feedstock chemicals. The styrenes themselves are useful as prepolymers, and are readily converted into other products by microorganisms.

5. Conversions of ferulic acid to vanillin: As a noteworthy goal, the conversion of ferulic acid to vanillin is of considerable interest. Some 12000 tons of synthetic vanillin are produced for flavoring purposes each year [57]. A number of bioconversions of ferulate to vanillin have been reported, but none of these accomplish the side-chain cleavage reaction (Figure 2 and Table 1) to yield the aldehyde (**5**) in good yield. Ishikawa *et al* [71] found that white rot fungi *Polyporus versicolor* and *Fomes fomentarius* could degrade ferulic acid to vanillin (**5**), which could be reversibly reduced to vanillyl alcohol (**6**) or oxidized to vanillic acid (**7**). Ötük found that *E. coli* metabolized ferulic acid to vanillin, then vanillic acid and protocatechuic acid [103]. Labuda *et al* [85] found that *Corynebacterium glutamicum* produced a mixture of vanillin (10–15% yield) and vanillic acid from ferulic acid. Iwahara *et al* [73] found an unidentified bacterial strain that converted ferulic acid to vanillic acid. Labuda *et al* [84] found that the conversion of ferulic acid to vanillin by *C. glutamicum*, *Aspergillus niger*, and *Pseudomonas putida* could be increased 10 to 100-fold by the addition of dithiothreitol in the medium. Addition of the thiol presumably precludes further reduction of the aldehyde functional group of vanillin by an aldehyde oxidoreductase.

6. O-Demethylation of ferulic acid to caffeic acid:

The demethylation of ferulic acid to caffeic acid (**3a**) is a common biotransformation reaction. Tillet and Walker [123] reported a *Penicillium rubrum* strain that O-demethylated ferulic acid to caffeic acid (**3a**, Figure 2). The caffeic acid thus formed is metabolized by conversion to protocatechuic acid, followed by further aromatic ring cleavage. Angeles de la Torre and Gomez-Alarcon found that anaerobic microorganisms from marsh sediment converted ferulic acid to caffeic acid [9]. Grbic-Galic found that the facultative anaerobe *Enterobacter* DG-6 was capable of O-demethylating ferulic acid to caffeic acid (**3a**, Figure 2) under either anaerobic or aerobic conditions. Anaerobically, caffeic acid was metabolized to cinnamic acid by ring dehydroxylation. Cinnamic acid was then reduced to phenylpropionic acid which was subsequently metabolized to phenylacetic acid. Aerobically, caffeic acid was oxidized to protocatechuic acid [48]. The mechanism of ferulate demethylation was proposed to involve a *trans*-methylation reaction resulting in the production of both **3a** and methanol [50]. Grbic-Galic and Young described a methanogenic

consortium that metabolized ferulic acid to caffeic acid, and then to CH₄ and CO₂ [51]. Scheline found that rat intestinal microflora could convert ferulic acid to *m*-hydroxyphenyl propionic acid, dihydrocaffeic acid and 4-ethylcatechol [114].

7. Reduction of ferulic acid to dihydroferulic acid (DFA):

Side-chain reduction of ferulic acid to dihydroferulic acid (**13**, Figure 2) has been observed frequently. The reduction is catalyzed by anaerobic microbial consortia [9,96], methanogenic consortia [51], *Pseudomonas cepacia* [8] and *Wolinella succinogens* [101] grown anaerobically, and by cultures of *Corynebacterium glutamicum* [85]. The side-chain reduction reaction also occurs in cultures of rat intestinal microflora [114] to give products including *m*-hydroxy-phenylpropionic acid, hydrocaffeic acid and 4-ethyl catechol [114]. Additional biotransformation reactions including dehydroxylation, demethylation, side-chain cleavage and further ring degradation often accompany the side-chain reduction reaction. Interestingly, the cinnamoyl side-chain reduction reaction of **1** to **13** was observed in our cultures of *Saccharomyces cerevisiae* [68]. With this yeast, **13** was obtained only when cells were incubated in an argon atmosphere. Under aerobic conditions ferulic acid was transformed to **14** (Figure 4). We suggested that the side-chain reduction reaction likely occurred by hydride attack of a quinoid intermediate (**25**) to afford **13** by an isomerization mechanism analogous to decarboxylations of ferulic acid and vanillic acid.

8. Biotransformations of vinylguaiacol:

Since 4-hydroxy-3-methoxystyrene (**14**) is a valuable bioconversion product from the extremely abundant ferulic acid, we examined microorganisms for their abilities to catalyze further transformations of **14**. Metabolic transformation studies of styrene by microorganisms have revealed several common pathways. Styrene is metabolized via styrene *cis*-glycol to 3-vinylcatechol and 2-vinylmuconate, demonstrating double bond and aromatic ring oxygenation prior to ring fission [131]. Metabolism of styrene via styrene oxide gives phenylacetaldehyde and phenylacetic acid [59,60,81]. With anaerobic consortia, after water addition to the double bond, styrene is oxidized through the phenylacetate route to yield a variety of products including ethylbenzene, phenylethanol, 1-phenylethanone, phenylacetaldehyde, phenylacetic acid, benzyl alcohol, benzaldehyde and benzoic acid [49]. Styrene transformations have also been reported to occur with *Pseudomonas putida* [17] and the black yeast *Exophala jeanselmei* [31].

Of some fifty microorganisms screened for their abilities to transform vinylguaiacol (**14**) we discovered a *Nocardia* strain that efficiently converted the substrate to products. When incubations were conducted under an argon atmosphere with resting cells of *Nocardia* spp, **14** was transformed to 4-ethyl-2-methoxyphenol (**28**, Figure 4) in 35% yield, 1-(4'-hydroxy-3'-methoxyphenyl)-ethanol (**29**) in 43% yield, and 1-(4'-hydroxy-3'-methoxyphenyl)-ethanone (**15**) in 2% yield (unpublished results). H₂¹⁸O was used as before in mechanistic studies to demonstrate (by MS analysis) that bioconversion of **14** to **29** involved the addition of water to the vinyl group of vinylguaiacol. We

speculated that a tautomerized intermediate such as **27** was involved in the transformation reaction. Deuterium oxide (D_2O)-containing media were used to demonstrate that the reaction sequence shown from **14** to **27** and **27** to **29** (Figure 4) were essentially reversible reactions. Recoveries of all products and determinations of deuterium incorporations by MS and deuterium NMR revealed that recovered **14** contained deuterium at the terminal olefin position; that **28** contained deuterium at both positions on the reduced ethyl side chain; and that both **29** and **15** contained deuterium in the terminal methyl position (unpublished results).

9. Selected biotransformations of vanillic acid: Although the biotransformation of ferulic acid is the major theme of this review, a few interesting biotransformations are described for vanillic acid (**7**), a major ferulic acid metabolite. Three of these are summarized in Figure 5. *O*-Demethylation of **7** affords protocatechuic acid (**10**) [124]. With *P. acidovorans*, *O*-demethylation requires NADH, H^+ along with formaldehyde, Fe^{2+} and reduced glutathione [124]. Conversion of vanillic acid to protocatechuic acid followed by ring-degradation is a common biodegradative pathway in microorganisms [21,22,43,56,75,79,97,113,124]. Vanillic acid may be degraded to quinone and acetophenone [75], and protocatechuic acid is degraded to 2-pyrone-4,6-dicarboxylic acid which is further metabolized to pyruvate and oxaloacetate [113] or β -keto adipic acid [97] after *ortho*-ring fission.

Pometto, Sutherland and Crawford found that *Strepto-*

myces setonii converted ferulic acid to vanillin, then vanillic acid. Vanillic acid was subsequently oxidized to protocatechuic acid and catechol with ultimate cleavage of the catechol aromatic ring to *cis,cis*-muconic acid [105]. Unspecified degradation products were obtained when protocatechuic acid was cleaved by a dioxygenase from *Bacillus subtilis* [56]. Quantitative nonoxidative decarboxylation of vanillate (10 mM) to guaiacol occurred with succinate-grown cells of *Bacillus megaterium* over a 9-h period [32]. Thus, guaiacol is a major metabolic product obtainable from ferulic acid.

We explored the mechanism of nonoxidative decarboxylation of vanillic acid by the same approach used in probing the mechanism of ferulic acid decarboxylation [69]. We proposed that **7** could isomerize to **30** (Figure 5) during the course of the reaction. Tautomerization of **7** to **30** involves addition to the aromatic ring of a proton from water which should be retained in guaiacol (**31**) following decarboxylation. The tautomerization is enzyme-mediated; the decarboxylation reaction is a chemical event because **30** is a vinylogous β -ketoacid that would spontaneously decarboxylate once formed. Mass spectral analysis of **31** indicated that a single deuterium atom was incorporated into guaiacol (**31**) obtained from incubation mixtures containing 50% or 100% deuterium oxide. Proton NMR spectroscopy also indicated that the proton at position 4 of **31** had been replaced by deuterium. Thus, the mechanism for guaiacol formation is similar to that identified for ferulic acid decarboxylation in that both systems apparently involve quinoid and vinylogous β -keto acid intermediates prone to decarboxylation.

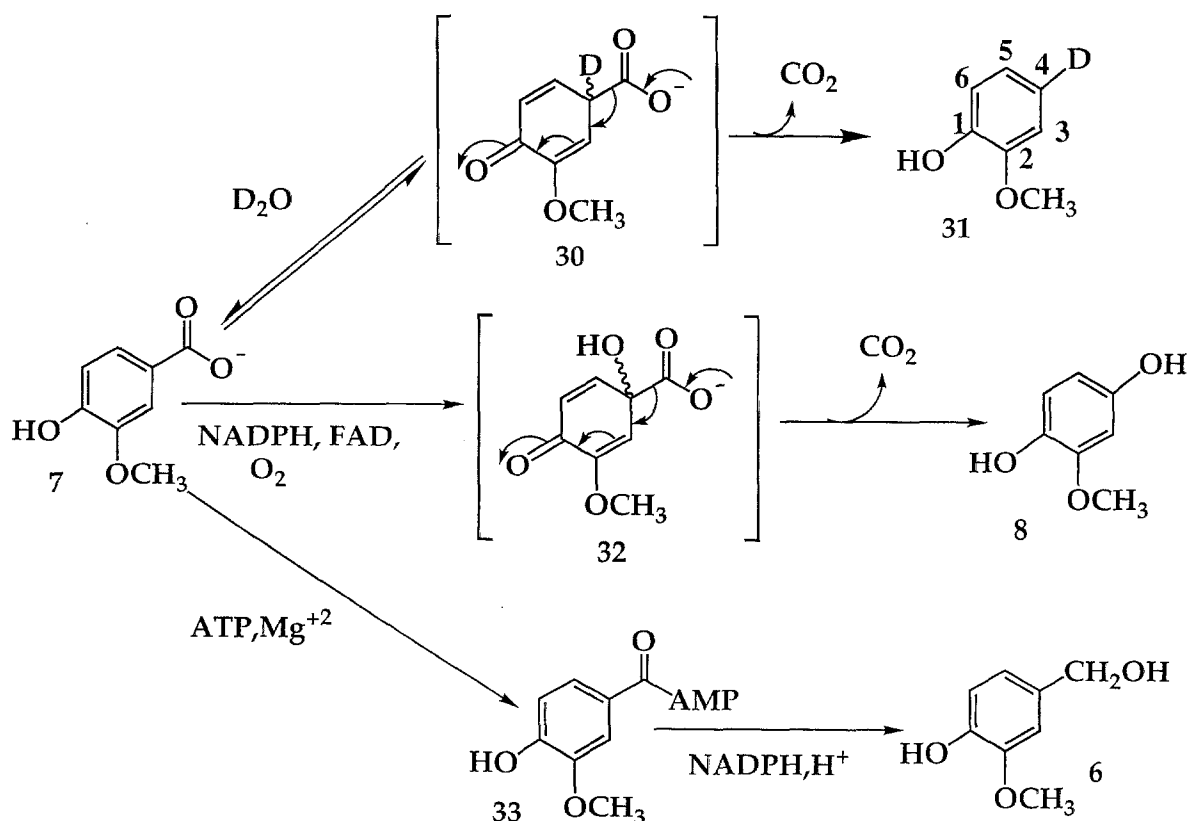


Figure 5 Selected oxidative and nonoxidative transformations of vanillic acid by microorganisms

Oxidative decarboxylation of vanillic acid to methoxyhydroquinone (**8**) (Figure 2) is also a common pathway for biotransformation in soft-rot, brown-rot and white-rot fungi [108]. Mycelial extracts prepared by sonication of cells of *Chaetomium*, *Humicola*, *Petriellidium*, *Daedalea*, *Lenzites*, *Fomes*, *Polyporus*, *Poria* and *Pycnoporus* contained high levels of an NADPH-dependent vanillate hydroxylase leading to the formation of **8**. Reaction progress was measured by the release of $^{14}\text{CO}_2$ from labeled vanillic acid. Vanillate decarboxylase from *Sporotrichum pulverulentum* requires both NADPH and molecular oxygen [20]. Vanillate decarboxylase from *S. pulverulentum* had a molecular weight of about 65 000 Da, had highest activity with both FAD and NADPH, required a *para*-hydroxyl group for decarboxylation, and worked best for vanillate, protocatechuate, *p*-hydroxybenzoate and 3,4-dihydroxybenzoate [23]. We suggest here that the oxidative decarboxylation mechanism leading to methoxyhydroquinone (**8**) involves a hydroxylated quinoid intermediate (**32**) that spontaneously decarboxylates to **8**.

Carboxylic acid reductases are a relatively unexplored group of enzymes. In many fungi and bacteria, carboxylic acid substrates like vanillic acid are activated by conversion to a carbonyl-AMP intermediate (Figure 5, **33**) followed by subsequent reduction to the benzylic alcohols [28]. The reaction is widespread in the microbial world, and we have observed its range to include aliphatic as well as aromatic acid substrates [28]. Facile reduction of carboxylic acids to their respective alcohols bears all of the interesting possi-

bilities of regio- and enantioselectivities with multifunctional substrates. Furthermore, this reaction is biotechnologically interesting because the carboxylic acid substrates are water soluble.

10. Oxidations of ferulates by peroxidases: A number of investigations have focused attention on enzymatic oxidations of ferulic acid or its derivatives. Many of these studies serve as a basis for modeling the formation of compounds in nature known as lignans or neolignans. The availability of enormous quantities of ferulic acid suggests its potential for use in the generation of oxygenated and biodegradable polymers by enzyme catalysis. The presence of the phenolic functional group obviously directs attention to the involvement of free radical oxidations for ferulic acid and its derivatives. Generation of phenolic radicals can theoretically lead to a plethora of dimeric, oligomeric or polymeric products. The aromatic ring and the conjugated double bond of the side-chain presents additional sites where electrons generated enzymatically or chemically can be delocalized. Examples of the formation of 'dimers' of ferulates formed naturally or 'in vitro' are common. Less well defined are the structures of polymers formed from ferulic acid and its relatives.

Most attention has been directed towards the action of peroxidases as biocatalysts. Chemical oxidants such as ferric chloride or ammonium persulfate, and photochemical oxidations have been employed to transform ferulates and to mimic the catalytic activities of peroxidases in nature.

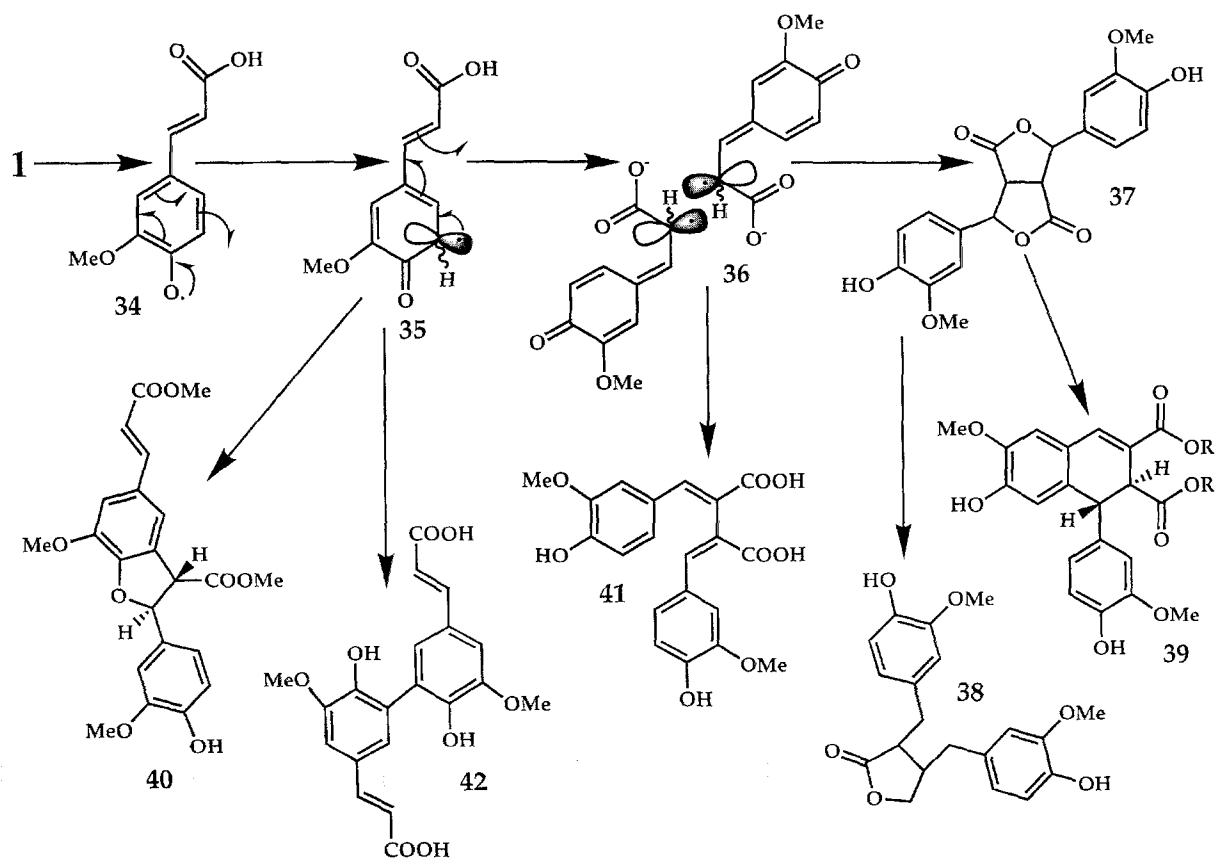


Figure 6 Products formed during enzymatic and chemical free radical oxidations of ferulic acid and derivatives

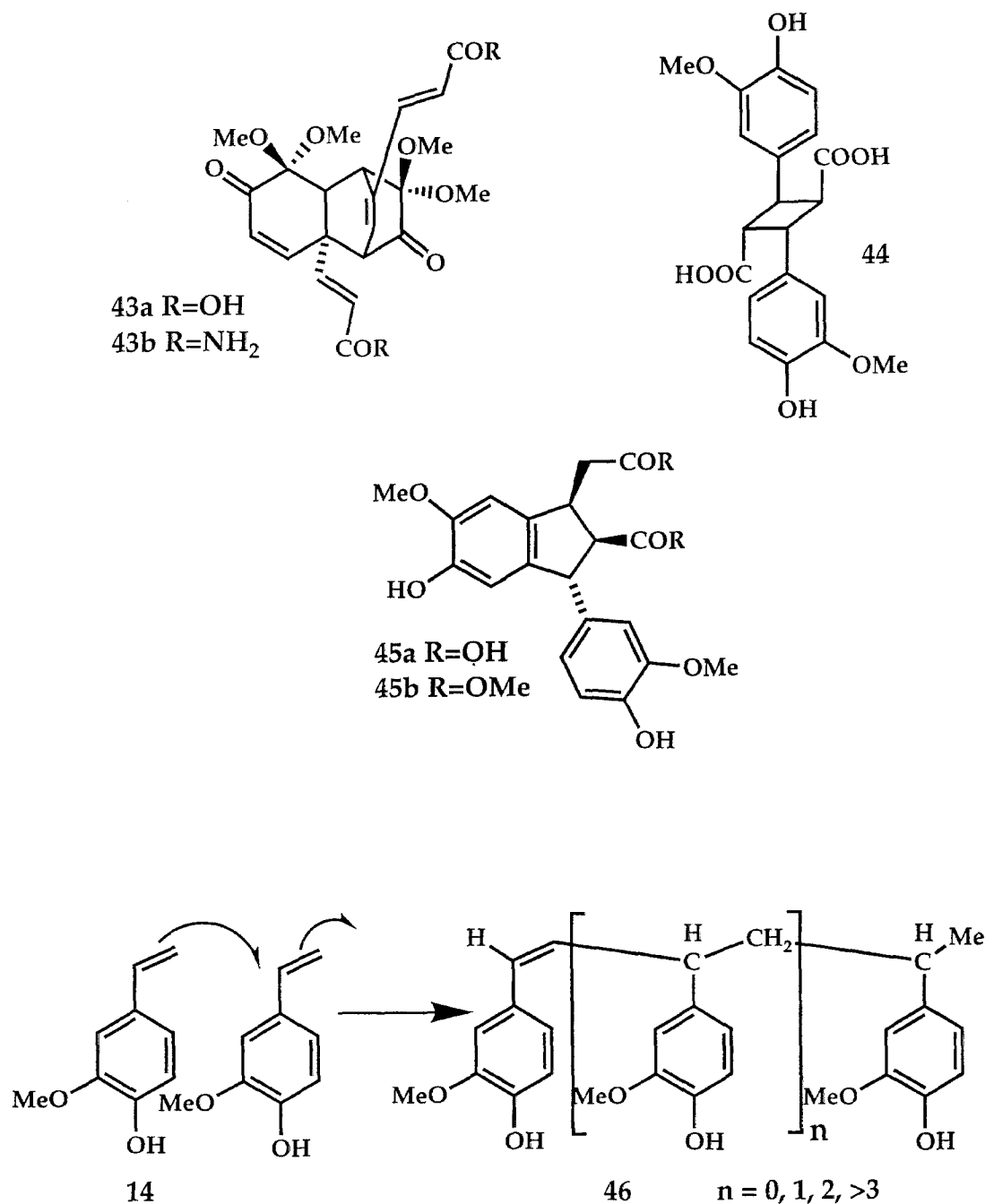


Figure 7 Structures of aryl indanes and some higher polymers of ferulic acid

Such work has a significant history. In 1944, Cartwright and Haworth studied the oxidation of ferulic acid with ferric chloride or ammonium persulfate [24]. These one-electron oxidants afforded the dilactone **37** as the major product (20% yield). Takei *et al* [121] demonstrated that **37** could be transformed into a matairesinol-like compound **38** by hydrogenolysis over palladium on carbon. Acidic treatment of **37** transformed the compound into the dihydronaphthalene derivatives **39** [121]. Thus, a pathway was established for the formation of **37** as an initial coupling product that serves as an intermediate for the formation of other lignan products. The initial steps (Figure 6) likely result from formation of phenolic radical (**34**) that is delocalized to the aromatic ring (**35**) and subsequently into the side-

chain as in **36**. Pairing of radicals in **36** followed by Michael-type addition ring closures gives **37**.

Oxidation of ferulic acid with horseradish peroxidase (HRP) gives complex mixtures of various products, many of which have not been characterized [15]. Ferulate methyl ester is transformed into the phenyl-coumaran type compound **40** in 25% yield with HRP [14,29]. This compound is also formed by oxidation with iodobenzene in the presence of tetraphenylporphyrinatomanganese (III) acetate (22%) [14,29] or with silver (I) oxide (44%) [11]. Ferulic acid has been used as a model substrate in studies with other peroxidases [45,128,135]. Different peroxidases appear to catalyze dimerization reactions in different ways. Caffeate peroxidase is reported to convert ferulic acid into

$\beta\beta'$ linked-type dimers such as **41** [45], while peanut peroxidase converts **1** into biphenyl type compounds like **42** [128]. Both compounds are formed by the initial oxidation of the phenolic functional group of **1** followed by electron delocalization to the ring as in **35** to form **42**, or to the unsaturated side-chain (**36**) to form **41**.

Electrochemical oxidations of ferulic acid afforded the asatone type dimer **43a** and the dilactone **37** as a minor product with yields depending on oxidation conditions [100]. Ferulamide is transformed only into the asatone type dimer **43b** in 56% yield electrochemically. Photochemical oxidation of ferulic acid by irradiation under 'daylight' fluorescent tubes affords low yields (*ca* 2%) of the truxilic acid-like compound **44** [44]. Photooxidation of its methyl ester with visible light in the presence of methylene blue as a photosensitizer afforded dihydronaphthalene **39** and phenyl coumaran **40** (less than 10% each) [83]. Ferulic acid and its methyl ester are also dimerized by standing in trifluoroacetic acid into aryl indane compounds **43a** and **43b** in 80% yield [4] (Figure 7). Similar products have not yet been reported by enzyme oxidation.

Higher polymers of ferulic acid were obtained by polymerization with horseradish peroxidase under weakly basic conditions (pH = 8) [112] or chemically with thionyl chloride [39] or oxalyl chloride [104]. Although the structures of the polymers remain undefined, these synthetic polymerized lignins demonstrated some level of anti-HIV and anti-tumor activities [13,82,95]. In studies on lignification in the cell wall to determine the nature of lignin-hydroxycinnamic acids interactions, some codimers from methyl 5-*O*-(*E*)-feruloyl- α -L-arabinofuranoside and conyferyl alcohol were synthesized [109,110].

Studies on the oxidative coupling of 4-hydroxy-3-methoxystyrene (**14**) obtainable enzymatically from ferulic acid have not been reported. However, the tendency for **14** to polymerize was confirmed by Kodaira *et al* who identified oligomers like **46** when the phenolic styrene was held at room temperature under an inert atmosphere in the dark [62,80]. A polymeric form of **14** was obtained by bulk polymerization in the presence of the chemical oxidant 2,2'-azobisisobutyronitrile as an initiator. The polymer was decomposed by microorganisms in soil, and four chemical degradation products were identified. Vanillic acid is formed, and ring cleavage of vanillic acid gives the monomethyl ester of β -carboxymuconic acid. These studies indicate that potentially useful and biodegradable polymers of phenolic styrenes can be generated readily by chemical and, yet to be demonstrated, enzymatic catalysis.

Conclusions

This review has focused attention on microbial and enzymatic transformations of ferulic acid. Related 'lignin' monomeric compounds *p*-coumaric acid and caffeic acid are also abundant in nature. Together, these phenylpropanoids are attractive preformed aromatic substrates of great value as precursors for other useful chemical products. The field of biocatalysis with these cinnamic acid derivatives is well established, and it is, in fact, quite old. Nevertheless, the exploitation of biocatalytic methods for the conversion of these abundant feedstock chemicals into other useful

products remains in its infancy. Many biotransforming enzymes remain to be isolated and characterized. As described in this review, ferulic acid has been transformed into useful phenolic aromatic chemical compounds of value as flavors (vanillin), leuco dyes for use in diagnostics (guaiacol), biodegradable polymer feedstock chemicals (vinylguaiacol), and dimeric products of potential value as drugs. Much work remains to be done. Preliminary investigations on mechanistic aspects of some enzymatic reactions have been reported, especially those dealing with conversions of ferulic acid to vanillic acid and vinyl guaiacol, and on the decarboxylation of vanillic acid into guaiacol. However, the catalytic mechanisms by which ferulate decarboxylase and vanillate decarboxylase function without apparent cofactor needs remain unknown. Although preliminary N-terminal sequences of some enzymes have been determined, none of the enzymes involved in reported biotransformations have yet been cloned and overexpressed. These approaches should yield an abundance of fruitful information about enzyme mechanism, and should avail sufficient amounts of enzymes for practical bioreactor applications.

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