

Utilization of dimeric lignin model compounds by mixed bacterial cultures

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Summary. The degradation of dimeric phenylpropanoid lignin model compounds using mixed bacterial cultures was studied. The six model compounds contained the most common linkages of lignin: β -O-4, β - β , β -5, and β -1. The results indicate that it is possible to enrich bacteria which are able to degrade all these compounds. Bacteria were also able to use these dimers as the sole source of carbon for growth. In view of these results it seems probable that bacterial inability to degrade polymeric lignin is due to the physical properties such as the molecular size of lignin.

Introduction

Fungi are believed to be mainly responsible for returning of lignin into the carbon cycle while bacteria may be important for the degradation of small fragments arising from fungal degradation of lignin (Ander and Eriksson 1978; Crawford 1981). Several authors have reported partial degradation by bacteria of synthetic lignin models (DHP) (Trojanowski et al. 1977; Haider et al. 1978; Kaplan et al. 1980), in situ ¹⁴C-phenylalanine labeled lignin of poplar (Odier et al. 1981), douglas fir (Crawford 1978; Crawford and Sutherland 1979, 1980; Phelan et al. 1979) or spruce (Robinson and Crawford 1978) or ¹⁴C-labeled lignosulfonic acids (Haider and Trojanowski 1981).

At least two reasons could explain why bacteria degrade lignin only partially. Firstly, the permeability barrier of the bacterial cell membrane may exclude molecules of the size of lignin. Metabolism could hence occur only if effectuated by extracellular catalysts. Secondly, lignin may contain chemical bonds which cannot be cleaved by bacteria. Before

taking on a study on the degradation of oligomeric lignin-born materials (industrial lignins) we studied the abilities of 11 mixed enrichment cultures of bacteria to utilize different types of dimeric phenylpropanoid model compounds. Degradation of the model compounds was determined by high performance liquid chromatography. The results are reported in this paper.

Materials and methods

Model compounds. The structures of compounds are presented in Fig. 1. 1-(3-methoxy-4-hydroxyphenyl)-2-(2-methoxyphenoxy)propane-1,3-diol (guaiacylglycerol- β -guaiacyl ether, compound I), was synthesized using the procedure described by Miksche et al. (1966). 1-(3,4-dimethoxyphenyl)-2-(2-methoxyphenoxy)propane-1,3-diol (veratrylglycerol- β -guaiacyl ether, compound II) was a gift from K. Lundquist, Chalmers University of Technology, Gothenburg. 1,2-bis(3-methoxy-4-hydroxyphenyl)-1,3-propanediol (compound III) was synthesized using a procedure described elsewhere (Ahvonen et al. 1983; Brunow et al. 1981). *d*-pinoresinol (compound IV) was from the same source as compound II and

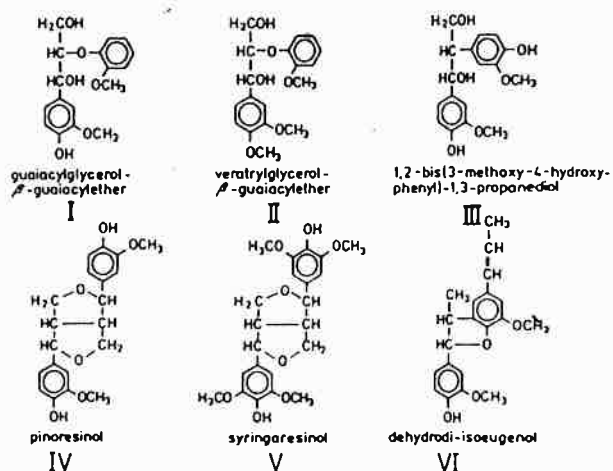


Fig. 1. The dimeric phenylpropanoid compounds used in the present study

d,l-syringaresinol (compound V) from T. Higuchi, synthesized by A. Kamaya, Institute of Wood Research, University of Kyoto. Dehydrodiisoeugenol (compound VI) was prepared from *trans*-isoeugenol according to Leopold (1950). The chemical identity and purity of each compound was verified with nuclear magnetic spectrometry and high performance liquid chromatography.

Cultures. The 11 mixed cultures used in this study originated from inoculae collected in an area which was polluted by pulp and paper industry waste waters. The inoculae were enriched for 6 months at room temperature on ten different media composed of liquors from pulping industry: thermomechanical pulping waste water (1); E₁-stage spent liquor of oxygen bleaching (2); black liquor at pH 9 (3) and 7.5 (4); O₂-stage effluent of oxygen bleaching with bark chips at pH 9.5 (5), at pH 9.5 with NH₄Cl (0.8 g/l), K₂HPO₄ (1.2 g/l), KH₂PO₄ (0.8 g/l), NaCl (0.04 g/l), and MgSO₄ (0.1 g/l) (6), and at pH 9.5 without addition of nutrients (7). The oxygen bleaching liquor was ultrafiltered with a filter with a cut-off of 5,000 D and the permeate and retentate were used in enrichments 8 and 9, respectively, at pH 8. Culture 10 was obtained from a bioreactor described elsewhere (Salkinoja-Salonen et al. 1983). Culture number 11 was forest soil without further enrichment. Waste waters from thermomechanical pulping contained 3.5–10 mg N/l and 0.16–23 mg P/l, bleaching liquors 6–8 mg N/l and 1.6–2.6 mg P/l, and black liquor ca. 14 mg N/l and ca. 2 mg P/l.

Dimer degradation experiments. The cultures were tested in the following way for their ability to utilize dimeric lignin model compounds as the sole source of carbon. A 50 µl inoculum of a mixed culture was diluted in 1 ml of a mineral salts medium containing (NH₄)₂SO₄ (1.0 mg), KH₂PO₄ (5.0 mg), MgSO₄ · 7 H₂O (0.1 mg), FeSO₄ · 7 H₂O (0.01 mg) per milliliter, supplemented with 5 µl of a vitamin mixture (Sundman 1964), 1.0 µl of a trace element mixture (Bauchop and Elsdon 1960) and 0.3 mg of compound I per milliliter of medium. The pH of the medium was adjusted to 7.0. The cultures were incubated aerobically without agitation in 5-ml tubes at 28° C and diluted after 7 days into a fresh medium (50 µl of culture into 1 ml of the same medium). These cultures were used as inoculae for experiments with the dimeric phenylpropanoid compounds I–VI (Fig. 1).

The effects of the antibiotics chloramphenicol and cycloheximide (Sigma, St. Louis, USA) (0.10 mg/ml) were examined using 1 ml of the liquid medium as above with 0.15 mg/ml of compound I.

An uninoculated control sample was prepared for each test incubation.

Measurement of bacterial growth. Bacterial growth was estimated by the increase in turbidity at 420 nm after settlement of insoluble substrate and possible precipitates. Biomass was calculated from the turbidity as described in the literature (Koch 1981). The content of protein of the cultures grown on dimers could not be determined by the Lowry procedure because of interference by phenolic substrates and the quantity of biomass protein was too small for other methods of protein assay.

Chromatography. At the end of the incubation (7–12 days) 1 ml of acetonitrile was added per milliliter of culture to dissolve unused substrate and possible reaction products. Insoluble solids (bacterial debris) were then removed by centrifugation. The acetonitrile extract was analyzed with a high performance liquid chromatograph (Micromeritics Instrument Corp., Norcross, Ga., USA) connected to a variable wavelength uv/vis-detector. Rad-Pak C18 (5 or 10 µm) columns were used in an RCM-100 compression module (Waters Associates Inc., Milford, Ma., USA). Acetonitrile and water mixtures containing 1% acetic were used as mobile phases.

Results

Tables 1 and 2 show how each of the cultures utilized the different model compounds. The percentage of degradation was calculated from the difference in the

Table 1. Degradation of model compounds I–VI^a

Culture	Compound					
	I	II	III	IV	V	VI
	Percent degraded of input					
1	100	17	81	100	42	0
2	100	100	49	24	17	4
3	100	100	94	80	62	0
4	100	100	91	100	30	6
5	100	100	96	0	47	34
6	100	100	86	100	66	78
7	100	100	79	100	44	0
8	100	24	22	95	39	9
9	100	53	20	0	38	42
10	100	100	95	100	37	2
11	62	6	0	8	–	–
Average degraded	97	73	65	64	42	18
Input of dimer (mg/ml)	0.3	0.3	0.5	0.3	0.5	0.5
Incubation time (days)	12	12	7	7	7	7

^a Degradation is expressed as $[(m_0 - m_x)/m_0] \times 100\%$, where m_0 = amount of dimer recovered from the uninoculated control sample, m_x = remaining amount of dimer at the end of the incubation

– = not determined

Table 2. Biomass yields for the cultures grown on lignin model dimers

Culture	Compound				
	I	II	III	IV	V
	Biomass (µg/ml dry weight)				
1	52.6	2.5	77.2	38.1	67.2
2	87.7	35.5	12.5	0.0	51.0
3	126.4	73.8	118.3	7.7	70.6
4	96.0	50.4	39.2	46.8	47.7
5	–	35.2	52.8	4.5	80.2
6	–	76.7	32.9	49.1	133.2
7	79.0	46.5	35.1	41.5	81.7
8	52.0	10.1	22.6	30.8	47.7
9	–	17.8	7.9	2.1	38.3
10	86.0	79.0	121.2	60.5	61.6
11	–	14.0	–	27.5	0.0
Average	82.8	40.1	52.0	28.1	61.7

Biomass yields were calculated from culture turbidities (A_{420}) (measured against uninoculated blank) according to Koch (1981)

– = not determined

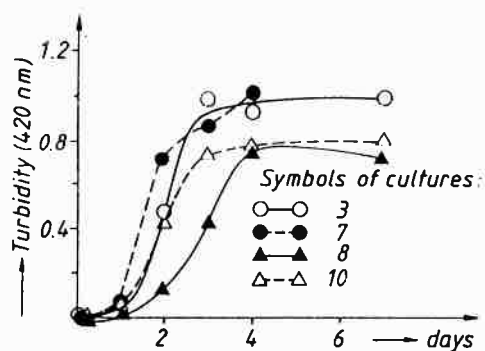


Fig. 2. Increase of culture turbidity during growth of cultures 3, 7, 8, and 10 on compound I (0.3 mg/ml)

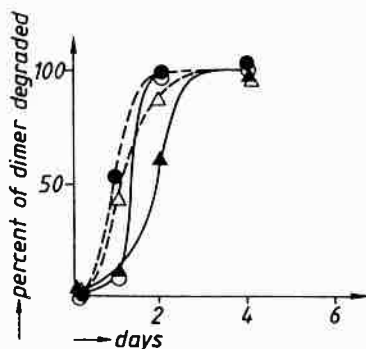


Fig. 3. Degradation of compound I by cultures 3, 7, 8, and 10. Symbols as in Fig. 2

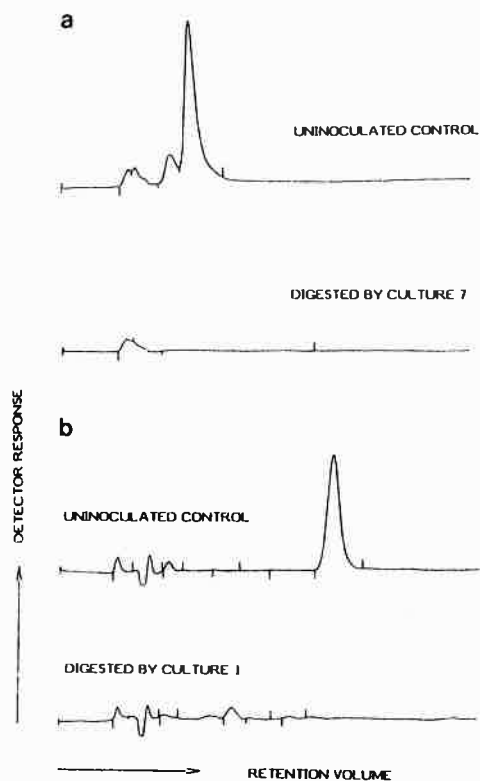


Fig. 4. Chromatographic analysis of the effect of bacterial digestion on compounds I (a) and IV (b)

amount of dimer in incubated samples with and without inoculum. Compounds with a β -O-4 ether bond (I and II) were well degraded by the cultures. This was expected because cultures pre-grown on compound I were used as inoculum and compounds I and II were closely related. However, the biomass yield was higher for compound I (average 82.8 μ g/ml) than for compound II (average 40.1 μ g/ml).

Figures 2 and 3 show that the maximum turbidity of the cultures was reached shortly after compound I had disappeared from the medium. The cultures grew to an optical density up to $A_{420} = 1.0$ (shown for compound I in Fig. 2). This is high considering the low substrate input (0.3 mg/ml). Many cultures grown on the other compounds (II–VI) were similarly dense.

The cultures grew turbid in a few days and in many cases no uv-absorbing (254 nm) compounds were left in the fully grown cultures, indicating disruption of aromatic rings. Examples of liquid chromatograms showing disappearance of the substrates are presented in Fig. 4.

Light microscopic and electron microscopic examination of the cultures showed that they were composed of bacteria exclusively. Table 3 shows how two antibiotics affected degradation of compound I. Chloramphenicol inhibited the degradation whereas cycloheximide had no effect. This confirmed the involvement of bacteria in the degradation; cycloheximide is known to affect eukaryotic cells and chloramphenicol prokaryotic cells.

The chromatograms showed that intermediate products were formed from compound III in several cultures. Two examples are presented in Fig. 5. Compound III used was pure *erythro* isomer. During incubation isomerization to *threo* isomer may have occurred. The peak area of the unincubated sample

Table 3. The effect of chloramphenicol and cycloheximide on the degradation of 1-(3-methoxy-4-hydroxyphenyl)-2-(2-methoxyphenoxy)-propane-1,3-diol (compound I) by cultures 3, 4, 7, and 10

Culture	Percentage of degradation ^a in the presence of		
	No antibiotic	Chloramphenicol ^b	Cycloheximide ^b
3	100	32	100
4	100	57	100
7	100	41	100
10	100	26	100
Uninoculated control	0	0	0

^a Calculated as in the Table 1. Initial concentration of compound I was 0.15 mg/ml and incubation time 4 days

^b Concentration 0.10 mg/ml

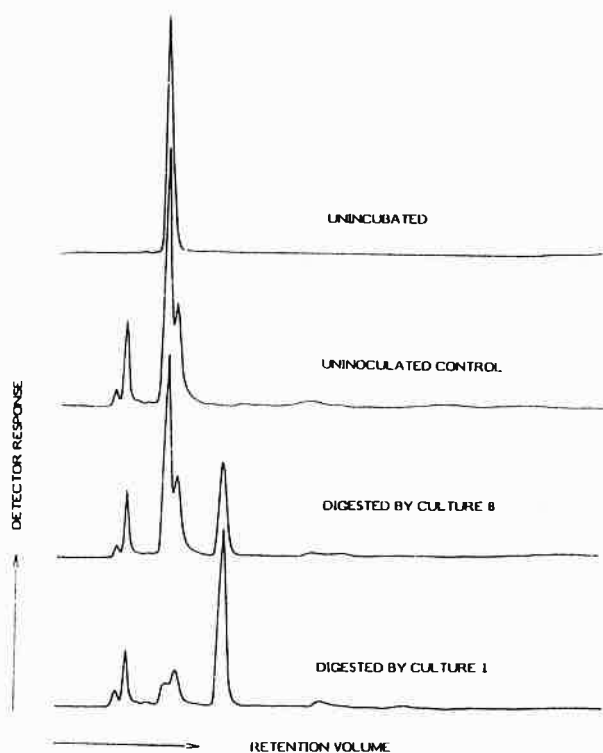


Fig. 5. Chromatographic analysis of products from compound III after digestion by bacteria

the combined area of the incubated control sample with the shoulder. The ratio of the peak areas (main peak : shoulder) in incubated control sample was 2 : 1.

Table 2 shows that biomass yield for compound III averaged 52.0 $\mu\text{g/ml}$. The average biomass yields for compounds IV and V were, 28.1 and 61.7 $\mu\text{g/ml}$, respectively, despite their similar structures.

Compound V was metabolized somewhat less efficiently (average 42%) than compound IV (average 64%). A pink colour emerged from compound V after a few days' incubation in some of the cultures. This may have been caused by accumulation of an intermediate product with quinonic structure (absorption maximum at 522 nm). Degradation of the compound V may have halted because of toxicity of such a quinone to the bacteria.

Only three of the cultures degraded compound VI to a significant degree. The biomass values were not included in Table 2 because biomass assay was disturbed by the insoluble substrate.

None of the compounds was completely resistant against bacterial attack. Culture 6 metabolized each of the six model compounds and produced biomass using these compounds as the sole source of carbon and energy. This culture was enriched by using oxygen bleaching waste water at pH 9.5 with addition of mineral salts.

Discussion

The most common linkages between the phenylpropanoid units of lignin are β -O-4-aryl ether, β -1, β - β , β -5, and 5-5 bonds (Lai and Sarkanen 1971). One way to study the biodegradability of lignin, is to use phenylpropanoid dimers linked to each other with one of these bonds. Bacterial degradation of such lignin models has been mostly studied with dimers containing the β -O-4 ether linkage (Trojanowski et al. 1970; Crawford et al. 1973; Fukuzumi and Katayama 1977; Katayama and Fukuzumi 1979b; Rast et al. 1980; Odier et al. 1982). Compounds of α,β -diaryl, pinoresinol and phenylcoumaran types have also been studied (Katayama and Fukuzumi 1978, 1979a).

The results presented here show that the independently prepared cultures selected for growth on one type of dimer [guaiacylglycerol- β -guaiacyl ether (I)] also utilized other types of dimers. Dehydrodiisoeugenol (VI) differed from the other phenylpropanoid dimers in that it was much less degraded.

Inspection of the cultures by light microscopy and electron microscopy showed that they were composed of bacteria. Some occasional fungal hyphae in the primary cultures disappeared during continued culture with guaiacylglycerol- β -guaiacyl ether (compound I) as the carbon source. Absence of fungal growth probably reflects the inability of fungi to use ligninous materials as the sole source of carbon (Ander and Eriksson 1978; Kirk 1981).

Our results resemble those described by Fukuzumi and Katayama (1977) and Katayama and Fukuzumi (1978, 1979a, b) who had isolated *Pseudomonas putida* (strain FK2) on 5,5'-dehydrodivanillic acid and found that it was also capable of utilising several different structurally unrelated dimers. This coincidence may point at some type of genetic or metabolic linkage in the biodegradation of these dimeric lignin models.

The results (Table 2) show further that some cultures not only degraded the different phenylpropanoid dimers so that the original compound completely disappeared, but biomass was also formed. Although Katayama and Fukuzumi (1978, 1979a, b) did not specify growth yields, it is obvious from their data that their *Pseudomonas* strain could also use several dimers as the source of carbon and energy. According to Crawford et al. (1973) a *Pseudomonas* strain E-3 was able to use compound II as its only source of carbon and energy. A low biomass density (10^7 cells/ml) was reported by Odier et al. (1981) for bacteria grown on poplar lignin. In the other reports where the degradation of dimeric, oligomeric or polymeric lignin models was studied with bacteria, a supplementing carbon source was always used (Tro-

janowski et al. 1977; Haider et al. 1978; Robinson and Crawford 1978; Kaplan and Hartenstein 1980; Crawford et al. 1981), probably because no significant growth was otherwise obtained.

The inability of bacteria to propagate efficiently on polymeric lignin might therefore be caused by the molecular weight of lignin or inaccessibility of the substrate to extra cellular enzymes. If bacterial metabolism of benzenoid compounds is confined to intracellular mechanisms, large molecules which are not transported into the cell cannot be metabolized. The permeability barrier of gram negative bacteria has been shown to be around 600–1,300 daltons for carbohydrates, polyethylene glycols, and peptides depending on the species (Payne and Gilvarg 1968; Nikaido and Nakae 1980). The permeability for lignin has not been determined. However, treatments of bacterial cells with organic solvents which are known to increase the permeability of cell membrane do not necessarily lead to decreased enzyme activity (Fukui and Tanaka 1982). This opens perspectives towards a lignin-based biotechnical process, since polymeric lignin can be converted into soluble oligomeric material by several known procedures, such as auto-oxidation under pressure (Sarkanen 1975), catalytic hydrogenolysis (Harris et al. 1938), acidolysis (Lundquist and Lundgren 1972; Lundquist 1976), hydrolysis (Lora and Wayman 1980), electrochemical oxidation or reduction (Brewer et al. 1948; Chum and Osteryoung 1981) or steam hydrolysis (Nimz et al. 1983).

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