

Immobilized Inocula of White-Rot Fungi Accelerate both Detoxification and Organic Matter Transformation in Two-Phase Dry Olive-Mill Residue

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The potential use for agronomic purposes of dry olive-mill residue (DOR), solid waste from the olive oil two-phase extraction process, might be impaired by its phytotoxicity. Although fungal treatments can detoxify DOR, long times are required for these processes. The objective of this study was to assess whether the addition of immobilized fungal inocula to DOR might improve colonization rates, thus reducing the time necessary for its detoxification and bioconversion. Inocula of *Panus tigrinus* CBS 577.79 and *Phlebia* sp. DABAC 9 immobilized on either chopped maize stalks or polyurethane sponge (PS) led to higher removals of both phenols and phytotoxicity from DOR than free inocula after 4 weeks of incubation. Best dephenolization (85%) was with PS-immobilized *Phlebia* sp., the use of which reduced germinability inhibition of *Lepidium sativum* and *Lactuca sativa* by 80 and 71.4%, respectively. Regardless of the type of inoculant, a low degree of humification was obtained.

KEYWORDS: White-rot fungi; immobilized inocula; phenols; dry olive-mill residue; phytotoxicity; thermochemolysis

INTRODUCTION

Large amounts of both liquid and solid organic wastes are annually produced by the olive oil extraction process, the seasonality of which leads to dramatic accumulation over a restricted time period. Environmental problems, formerly confined within the Mediterranean area and due to both improper storage and disposal procedures, are rapidly evolving in other countries such as Australia, Chile, and the United States. At the beginning of the 1990s, an innovative two-phase extraction process (TPEP) was introduced in Spain and in Croatia, where this technology rapidly extended to about 90% of the olive oil mills (1). The TPEP generates a liquid phase (olive oil) and a water-rich solid organic waste (alpeorujo), which is dried and extracted with solvents to obtain an extra yield of oil and the dry olive-mill residue (DOR). It has been estimated that the annual production of DOR in Spain approaches 4 million tons (2). Among the possible strategies of the two-phase olive-mill wastes upgrading, much emphasis has been given to their agronomic use after composting (1). However, these wastes exhibit peculiar physical and chemical properties that hinder the success of the biological approach; both the high moisture content and the dough-like texture limit gaseous exchanges, thus slowing the process (3). The preponderance of plant cell wall macromolecules (i.e., lignin, cellulose, and hemicellulose) associated with both a relatively low amount of nitrogen (0.7-1.8%) and the presence of antimicrobial compounds makes its composting slower (from 28 to 40 weeks) than that of other lignocellulosic wastes (3, 4).

DOR has a high concentration of organic matter (OM), which also includes toxic compounds, capable of inhibiting microbial growth (5) and germination and vegetative growth in plants (6). Because DOR toxicity has been mainly ascribed to phenols (2, 7), the use of ligninolytic fungi capable of degrading such compounds (6) can be an adequate approach for upgrading the waste. White-rot fungi (WRF) have been shown to remove phenols from DOR, although detoxification required long colonization times (7).

For this reason, the main objective of this study was to assess whether the addition of immobilized inocula of WRF to DOR might improve the colonization rate and reduce times required to detoxify the waste. In the present study, both natural and synthetic carriers, namely, chopped maize stalks (CMS) and polyurethane sponge (PS), were used. Although use of the latter carrier cannot be envisaged at the field scale, PS was used in this study because, besides being a valuable immobilization support for fungi, it is totally inert, thus allowing possible comparative discrimination between protective and trophic effects.

Panus tigrinus CBS 577.79 and *Phlebia* sp. DABAC 9, previously selected on the basis of their colonization rate on agar media added with DOR, were inoculated on CMS or PS and 7-day-old colonized carriers subsequently added to DOR. A particular focus was given in this study to the determination of the impact of the selected fungi on the relative abundances of

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Article

both labile and stabilized carbon fractions in DOR, as opposed to the majority of investigations on the fungal conversion of this waste (6). In addition, possible structural changes in the waste that had undergone fungal treatment were investigated by pyrolysis with tetramethylammonium hydroxide followed by gas chromatography mass-spectrometry (TMAH-Py-GC MS) and compared with corresponding abiotic controls.

MATERIALS AND METHODS

Materials and Preparation of Immobilization Supports. DOR, withdrawn from an olive oil manufacturer (Sierra Sur S.A., Granada, Spain), was stored at -20 °C until used. Its chemical composition was as follows: pH, 5.13; total organic carbon, 49.5%; total nitrogen, 2.28%; total phosphorus, 0.21%; lignin, 24.7%; cellulose, 18%; hemicellulose, 12.8%; total phenols, 3.18%; total lipids, 0.2%; ashes, 9.2%. The most abundant elements in DOR were (g kg⁻¹) potassium, 30.5; calcium, 13.6; magnesium, 3.8; iron, 1.1; sodium, 0.17; copper, 0.07; zinc, 0.06; and manganese, 0.04. PS cubes, 0.5 cm width each, were rinsed with water in a 1:20 (w/v) ratio and autoclaved (121 °C for 20 min) twice prior to their use. Maize stalks were chopped into approximately 0.5 cm pieces and sieved. Hydroxytyrosol, oleuropein, and luteolin standards for chromatography were from Extrasynthese (Lyon, France), whereas the remaining standards were from Aldrich (Milan, Italy).

Organisms and Inoculum Preparation. *Panus tigrinus* (CBS 577.79) and *Phlebia* sp. (DABAC 9) were maintained at 4 °C and routinely subcultured every month on potato dextrose agar slants. Fungal inocula were prepared as reported elsewhere (8). Five milliliters of the inoculum (ca. 50 mg of dw) was aseptically added to 50 g of sterilized supports and incubated at 28 °C for 7 days.

Sample Preparation and Fungal Treatment. Deionized water was added to DOR to adjust the moisture content to 25% (w/w) prior to sterilization (120 °C for 20 min). Then, either free mycelium (65 mg of dw) or colonized solid supports (2.5 and 0.24 g of CMS and PS, respectively), were covered with 17.5 g of DOR. The amounts of either free or immobilized mycelial biomass to which the waste was added were normalized on the basis of the same initial chitin content (5 mg). Solid-state cultures on DOR were carried out at 28 °C in the dark under stationary conditions for 30 days. Non-inoculated and sterilized DOR samples were prepared and incubated as above and will be referred to as abiotic controls from here onward. In addition, biotic controls were prepared by mixing in a 10:1 ratio (w/w) sterilized DOR with a sandy-loamy soil, the physicochemical properties and microbial density of which were reported elsewhere (8). All experiments were carried out in triplicate.

Analytical Assays. Samples were incinerated at 600 °C for 12 h, and OM was obtained by subtracting ash content from the whole sample weight. OM losses (%) were calculated from the ash contents in the waste at start (X_1) and in either fungal-treated DOR or corresponding incubation controls (X_2) according to eq 1 (7):

OM losses (%) =
$$100 - \frac{[X_1 \times (100 - X_2)]}{[X_2 \times (100 - X_1)]} \times 100$$
 (1)

Total extractable carbon (TEC) was extracted by mechanical shaking for 48 h at 65 °C with 0.1 M NaOH plus 0.1 M Na₂P₄O₇ under N₂ atmosphere and using a 1:50 solid/liquid ratio. The suspension was then centrifuged (8000g; 20 min) and the supernatant filtered through a 0.8 μ m membrane (Millipore, Danvers, MA). Humic acid carbon (C_{HA}) was precipitated from the filtrate by adding H₂SO₄ up to pH 2.0, then centrifuging as above, and the pellet was collected and stored at 4 °C (9). The supernatant (25 mL) was loaded onto a small column (10 mL) packed with polyvinylpolypyrrolidone and previously equilibrated with 5 mM H₂SO₄; the retained fraction made of fulvic acid carbon (C_{FA}) was then eluted by 0.1 M NaOH and collected into a centrifuge tube (9). Total organic carbon (TOC), TEC, C_{HA}, and C_{FA} were determined according to the dichromate oxidation method as previously reported (7). The loss of C throughout incubation was calculated according to eq 2 (4)

TOC losses (%) =
$$100 - \frac{[X_1 C_2]}{[X_2 C_1]} \times 100$$
 (2)

where X_1 and X_2 and C_1 and C_2 are ash and TOC concentrations,

respectively, in the waste at start and in either abiotic controls or related fungal treatments. Degree of humification (DH), humification ratio (HR), and humification index (HI) were calculated from the following ratios, respectively:

$$DH\% = \frac{C_{HA} + C_{FA}}{TEC} \times 100$$
(3)

$$\mathrm{HR\%} = \frac{C_{\mathrm{HA}} + C_{\mathrm{FA}}}{\mathrm{TOC}} \times 100 \tag{4}$$

$$\mathrm{HI} = \frac{C_{\mathrm{HA}}}{\mathrm{TOC}} \times 100 \tag{5}$$

Total N, P, K, Fe, Mn, Cu, and Zn contents of the DOR were determined as reported elsewhere (7). Acetone/water-soluble phenols (AWSP) were extracted from 2 g of DOR with 20 mL of acetone/acidified water at pH 3.0 (50:50, v/v) for 24 h at 4 °C and determined using tyrosol as the standard (7). Dephenolization efficiency (DE) was calculated by normalizing the amount of phenols removed to the same fungal biomass (g of phenols removed g^{-1} of chitin). On the other hand, the dephenolization selectivity (DS) was calculated by the ratio of removed phenols to organic matter depleted.

Chromatographic and TMAH-Py-GC MS Analyses. Gas chromatographic—mass spectrometric (GC-MS) analyses were performed on DOR ethyl acetate residues derivatized with N,O-bis(trimethylsilyl) trifluoroacetamide in pyridine as previously reported (10). Identification of aromatic compounds was based on comparison with retention times and mass spectra of pure standards and comparison of the mass spectra with those in the NIST92 and NIST95 libraries.

Reversed-phase high-performance liquid chromatography (RP-HPLC) was used to quantify the degradation of the main aromatic compounds (11).

TMAH-Py-GC MS analyses were performed as previously described (12). Pyrolysis products were identified both by comparing mass spectra with data in the NIST02 library and by interpreting the fragmentation pattern. Data are the mean of triplicate runs, and the percentages of pyrolysis products were calculated from the relative areas of the peaks after recalculation according to the exact weight of samples. With the notable exception of pyrolysis products derived from carbohydrates, the coefficient of variation of the most abundant peaks was lower than 15%. Individual fragments detected after pyrolysis were divided into the following four main groups according to their origin: lignin-related compounds (L), carbohydrates (C), fatty acids (FA), and nitrogen-containing compounds (N). When possible, fragments belonging to the first group were assigned to lignin basic structures: P, p-hydroxyphenyl; G, guaiacyl (3-methoxy-4-hydroxyphenyl); and S, syringyl (3,5-dimethoxy-4hydroxyphenyl). The relative percent abundances of P, G, and S subunits were calculated by dividing the areas of peaks pertaining to each subunit either by the total area of pyrograms (P%, G%, S%) or by their sum (Pn%, Gn%, Sn%). The syringyl/guaiacyl ratio (S/G) and guaiacyl and syringyl acid/aldehyde ratios, namely, (Ad/Al)_G and (Ad/Al)_S, were calculated as described by Vane et al. (13).

Fungal Growth and Metabolic Activity. The extent of fungal biomass was indirectly estimated by the chemical determination of the chitin content in the solid substrate as previously described (7). Growth was also assessed subjectively on a 0-10 scale, with 0 meaning absence of growth and 10 complete hyphal colonization of the surface of the solid substrate (14). The index of metabolic activity in axenic fungal cultures, namely, the fungal metabolic index (FMI), was calculated from the mass ratio of organic matter losses to fungal biomass and expressed as grams of substrate consumed per gram of glucosamine.

Enzyme Assays. Extracellular enzymes were extracted from DOR, dialysis-filtered, and 20-fold concentrated as previously reported (8). Manganese peroxidase (MnP) and laccase activities were assayed as described by D'Annibale et al. (10). The monophenolase activity of tyrosinase, that is, the hydroxylation of monophenols to *o*-diphenols, was assayed according to the method of Espín et al. (15). Appropriate controls were performed with heat-denatured extracts. Enzyme activities were expressed as nanokatals per gram of DOR.

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Table 1. Fungal Growth Indices, Organic Matter (OM) and Total Organic Carbon (TOC) Losses, Fungal Metabolic Index (FMI) in DOR at Start (DOR *t*₀), Its Biotic Control (DOR-BC), and the Same Waste That Had Been Incubated with *P. tigrinus* and *Phlebia* sp. Added either in Free Form (DOR-*Pt* and DOR-*P*sp., Respectively) or Previously Immobilized onto Chopped Maize Stalks (DOR-*Pt*-CMS and DOR-*P*sp-CMS, Respectively) or Polyurethane Sponge (DOR-*Pt*-PS and DOR-*Ps*p-PS, Respectively) and in Abiotic Controls (AC-CMS and AC-PS)

sample	growth index				
	subjective ^a (0-10)	chitin ^a (mg/g)	OM loss ^a (%)	TOC loss ^a (%)	FMI ^a
DOR t ₀	nd ^b	0.2 A	nd	nd	nc ^c
DOR-BC	nd	0.4 B	7.0 B	8.5 C	nc
DOR-AC-CMS	nd	0.1 A	4.0 A	5.8 B	nc
DOR-AC-PS	nd	0.1 A	3.7 A	4.9 A	nc
DOR-Pt	3	0.7 C	8.3 B	10.5 D	118.3 B
DOR-Psp	4	0.7 C	7.2 B	8.4 C	109.9 B
DOR-Pt-CMS	8	1.5 D	24.3 E	24.1 G	158.7 C
DOR Psp-CMS	8	1.4 D	32.5 F	35.0 H	235.8 D
DOR-Pt-PS	8	1.4 D	16.9 D	19.5 F	117.2 B
DOR-Psp-PS	7	1.5 D	11.4 C	16.1 E	78.4 A

^a Data are the mean of three independent experiments: column means followed by the same upper case letter are not significantly different (*P* ≤ 0.05). ^b nd, not determined. ^c nc, not calculated.

Table 2. Acetone/Water-Soluble Phenols (AWSP) and Ethyl Acetate-Extractable Phenols (EAEP), Dephenolization Efficiency and Selectivity in DOR at Start (DOR t_0), Its Biotic Control (DOR-BC), and the Same Waste That Had Been Incubated with *P. tigrinus* and *Phlebia* sp. Added either in Free Form (DOR-*Pt* and DOR-*Psp*, Respectively) or Previously Immobilized onto Chopped Maize Stalks (*Pt*-CMS and *Psp*-CMS, Respectively) or Polyurethane Sponge (*Pt*-PS and *Psp*-PS, Respectively) and in the Respective Abiotic Controls (AC-CMS and AC-PS)

sample	$AWSP^{a} (mg g^{-1})$	$EAEP^{a} (mg g^{-1})$	dephenolization efficiency	dephenolization selectivity	
DOR to	41.9 D	11.7 E	nc ^b	nc	
DOR-BC	35.9 D	10.4 D	nc	nc	
DOR-AC-CMS	39.4 D	11.6 E	nc	nc	
DOR-AC-PS	37.4 D	12.0 E	nc	nc	
DOR-Pt	27.4 C	7.7 C	20.7 A	0.17 C	
DOR-Psp	23.8 B	6.7 C	27.9 C	0.25 E	
DOR-Pt-CMS	6.0 A	1.2 B	23.4 B	0.15 B	
DOR-Psp- CMS	5.3 A	0.6 A	26.5 BC	0.11 A	
DOR-Pt-PS	6.2 A	1.0 B	24.8 B	0.21 C	
DOR-Psp-PS	5.8 A	0.4 A	24.7 B	0.32 D	
				,	

^a Data are the mean of three independent experiments: column means followed by the same upper case letter are not significantly different ($P \le 0.05$). ^b nc, not calculated.

Phytotoxicity Assays. The phytotoxicity of DOR, at start or differently treated, was determined as described by Zucconi et al. (16). Seventy seeds of either *Lepidium sativum* or *Lactuca sativa* were placed, in triplicate, in 9 cm diameter Petri dishes containing sterilized quartz sand mixed with 5% (w/w) DOR and incubated in the dark at 25 °C for 48 h. Control experiments were conducted in the absence of the waste. Germinated seeds were counted and radicle growth measured. Percent inhibition (I%) of either germinability or radicle elongation was calculated from eq 6

$$I\% = \left(1 - \frac{G}{G_c}\right) \times 100 \tag{6}$$

where G is the number of either germinated seeds or radicle length observed in the presence of either fungal-treated DOR or abiotic controls and G_c the same parameters in the absence of the waste.

Statistical Analysis. Multiple pairwise comparisons were performed by the Tukey test. Thermochemolysis data were also subjected to principal component analysis (PCA) by the use of the Simca-P 8.0 software (Umetrics, Umea, Sweden). The possible presence of either moderate or strong outliers in observations was checked by the squared prediction errors and hotelling (T^2) of t scores, respectively (17). For both ANOVA and PCA, percent data were normalized by arcsin of the square root transformation.

RESULTS

Effect of the Carrier on both Mycelial Growth and Metabolic Activity on DOR. The white and yellowish mycelia of *P. tigrinus* and *Phlebia* sp., respectively, became visible at 2 weeks after inoculation but, at the end of the set incubation time, colonization of the substrate was not complete as assessed by both subjective growth index and chitin content (Table 1). By contrast, both fungi

added to DOR in immobilized form exhibited rapid growth on the waste, and colonization of the solid substrate was extensive after 4 weeks. The fungal growth was not significantly affected by the inoculum carrier; the chitin contents of the waste colonized by the two fungi were rather similar (**Table 1**).

Despite the large similarities in the amount of growth, the bioconversion capabilities of the two immobilized fungi were markedly affected by the type of carrier, as indicated by both OM and TOC losses. *Phlebia* sp. was more affected by the type of carrier, leading to the highest (32.5%) and lowest (11.4%) OM losses when immobilized onto CMS and PS, respectively (**Table 1**). Moreover, CMS proved to be a better support than PS for *P. tigrinus*, leading to higher extents of both OM and TOC losses ($24.3 \pm 1.2 \text{ vs} 16.9 \pm 0.5$, respectively, and $24.1 \pm 0.8 \text{ vs} 19.5 \pm 0.4$, respectively). The FMI values were similar in DOR inoculated with free mycelia of the two fungi. By contrast, *Phlebia* sp. was greatly affected by the inoculum carrier, resulting in the highest and lowest FMI values with CMS and PS (235.8 vs 78.4, respectively) (**Table 1**).

Dephenolization of DOR and Production of Extracellular Phenoloxidizing Enzymes. The AWSP content of the waste (i.e., 41.9 g kg^{-1} of DOR) did not change in both abiotic controls, whereas about 14% dephenolization was detected in the biotic control (**Table 2**). AWSP percent removals in DOR that had been incubated with free mycelia of *P. tigrinus* and *Phebia* sp. were 36 and 43%, respectively; by contrast, the extent of removal was significantly higher with both immobilized fungal inocula. In fact, AWSP were depleted from DOR by about 85% by both CMSand PS-immobilized *P. tigrinus*. Similar removals (i.e., 87%) were **Table 3.** Initial Concentrations of Ethyl Acetate-Extractable Aromatic Compounds and Percent Removals of Each Compound Observed in DOR That Had Been Colonized by *Panus tigrinus* or *Phlebia* sp. Added either in Free Form (NS) or Previously Immobilized onto Chopped Maize Stalks (CMS) or Polyurethane Sponge (PS)^a

		% removal ^b					
	initial concn (mmol/kg)	Panus tigrinus			Phlebia sp.		
compound		NS	CMS	PS	NS	CMS	PS
protocatechuic acid 1	0.2	55.1	95.0	97.0	58.9	97.7	100
hydroxytyrosol 2	18.0	29.6	93.6	94.9	37.7	98.2	99.5
catechol 3	3.8	62.0	93.1	88.9	70.3	90.7	93.6
tyrosol 4	6.9	36.2	80.9	96.4	48.7	85.0	89.4
3,4-dihydroxymandelic acid 5	1.8	60.0	100	100	68.0	100	100
vanillic acid 6	4.0	68.9	84.7	87.7	69.6	97.0	98.0
syringic acid 7	2.3	0.0	0.0	45.0	65.5	53.0	54.8
<i>p</i> -coumaric acid 8	1.4	47.0	35.0	14.3	53.0	91.1	96.8
<i>p</i> -methoxyphenylacetic acid 9	0.8	0.0	67.6	65.5	0.0	81.9	80.2
3,4,5-trimethoxybenzoic acid 10	0.6	0.0	42.3	89.9	0.0	55.5	62.0
luteolin 11	0.4	0.0	97.0	97.0	0.0	95.0	98.0

^a Arabic numbers have been assigned to each compound in order of increasing retention time ($t_{\rm R}$): **1**, 10.11 min; **2**, 10.6 min; **3**, 13.9 min; **4**, 15.5 min; **5**, 16.9 min; **6**, 17.6 min; **7**, 19.2 min; **8**, 24.0 min; **9**, 25.9 min; **10**, 29.9 min; **11**, 41.3 min. ^b Data are the mean of triplicate runs, and the relative standard deviation was \leq 8%.

attained with DOR that had been incubated with both CMS and PS-immobilized *Phlebia* sp. Despite the large differences between free and immobilized mycelia in the overall amounts of phenols removed, it was found that DE values were highest in DOR incubated with both free and CMS-immobilized *Phlebia* sp. (27.9 and 26.5 g of phenols removed mg⁻¹ of chitin, respectively). On the other hand, DS values showed that the highest and lowest selectivities were observed with PS- and CMS-immobilized *Phlebia* sp. (0.32 and 0.11, respectively) (**Table 2**).

Both GC-MS and reversed-phase HPLC analyses of ethyl acetate DOR extracts led to the identification of 11 aromatic compounds (**Table 3**). Compounds **2** and **4** were by far the most abundant aromatic components in ethyl acetate extracts (18 and 6.9 mmol kg⁻¹, respectively). However, significant concentrations of the substituted benzoic acids **1**, **6**, and **7** and the cinnamic acid derivative **8** were detected (0.2, 4.0, 2.3, and 1.4 mmol kg⁻¹, respectively).

Low laccase levels (0.2 nkatal g^{-1}) were detected in the biotic control (**Table 4**). In DOR that had been incubated with free mycelium of *Phlebia* sp., both laccase and MnP activities were found (2.1 and 4.6 nkatal g^{-1} , respectively), whereas in *P. tigrinus* cultures only the former was detected. In DOR incubated with PS-immobilized fungi, both laccase and monophenolase activities were higher than with CMS-immobilized fungi. Regardless of the type of carrier, MnP was detected only with PS-immobilized *Phlebia* sp. (46.3 nkatal g^{-1} DOR).

Fungal Removal of Phytotoxicity from DOR. Both germinability and radicle elongation of *Lactuca sativa* and *Lepidium sativum* were suppressed in the presence of 5% DOR, thus showing the high phytotoxicity of the waste. These toxic effects were not significantly removed in either abiotic and biotic controls of DOR (**Figure 1**) and were largely evident in the waste that had been inoculated with free mycelia of both fungi.

Phytotoxicity toward *Lepidium sativum* was generally attenuated in the waste that underwent incubation with immobilized fungal inocula. Best results were obtained with PS-immobilized *Phlebia* sp.: inhibition of germinability and radicle elongation amounted to 20 and 2.8%, respectively. By contrast, *P. tigrinus* was unable to mitigate the waste phytotoxicity toward *L. sativa*, regardless of the use of free or immobilized inocula. With this plant, the lowest inhibition of germinability (19.6%) was observed in the waste treated with CMS-immobilized *Phebia* sp. (Figure 1A).

Regardless of the treatment typology, germinability of both plants plotted versus the absolute amounts of added phenols

Table 4. Laccase, Mn-Dependent Peroxidase (MnP), and Monophenolase Activities in DOR at Start (DOR *t*₀), Its Biotic Control (DOR-BC), and the Same Waste That Had Been Incubated with *P. tigrinus* and *Phlebia* sp. Added either in Free Form (DOR-*Pt* and DOR-*Ps*p, Respectively) or Previously Immobilized onto Chopped Maize Stalks (*Pt*-CMS and *Psp*-CMS, Respectively) or Polyurethane Sponge (*Pt*-PS and *Psp*-PS, Respectively) and in the Relative Abiotic Controls (AC-CMS and AC-PS)

	enzy	enzyme activity ^a (nkatal g^{-1} of DOR)					
sample	laccase	MnP	monophenolase				
DOR t ₀	nd ^b	nd	nd				
DOR-BC	0.2 A	nd	nd				
DOR-AC-CMS	nd	nd	nd				
DOR-AC-PS	nd	nd	nd				
DOR-Pt	1.6 C	nd	nd				
DOR-Psp	2.1 C	4.6 A	nd				
DOR-Pt-CMS	7.7 D	nd	0.6 A				
DOR-Psp-CMS	1.0 B	nd	0.5 A				
DOR-Pt-PS	9.7 E	nd	0.8 B				
DOR-Psp-PS	2.3 C	46.3 B	0.7 B				

^{*a*} Data are the mean of three independent experiments: column means followed by the same upper case letter are not significantly different ($P \le 0.05$). ^{*b*} nd, not detected.

exhibited sigmoidal dose–response curves from which the inhibitory concentrations leading to 50% of germination inhibition (IC₅₀) were calculated (3.6 and 2.7 g kg⁻¹ soil for *Lepidium sativum* and *Lactuca sativa*, respectively) (**Figure 2**). The degree of deviation from the fitting curves was highest with free mycelium of *Phlebia* sp.

Effect of Fungal Growth on Structural Changes and Carbon Pools in DOR. On the basis of the limited impact of some treatments on both dephenolization and detoxification, investigations on possible structural changes and shifts in the relative abundances of carbon pools in DOR were limited to only abiotic controls and to experiments conducted with immobilized fungal inocula.

TMAH-Py-GC MS of DOR abiotic control yielded more than 50 fragments, most of which were methyl ethers and esters of carbohydrates, fatty acids, and aromatics. The majority of these fragments were identified, as shown in **Figure 3A**, and found to be aromatic compounds. Among them, 1,4-dimethoxybenzene and 1,2,4-trimethoxybenzene were dominant; these compounds, however, have been shown to derive from both lignin and cellulose thermochemolysis (*18*). On the other hand, the major aromatic fragments unambiguously derived from lignin were 3, 4-dimethoxybenzoic acid methyl ester (2.7%) and 3,4-dimethoxystyrene (1.6%). The relative abundances of lignin-related



Figure 1. Percent inhibition of germinability (**A**) and radicle elongation (**B**) in *Lepidium sativum* (black bars) and *Lactuca sativa* (white bars) seedlings grown at 27 °C for 48 h in quartz sand treated with 5% (w/w): DOR at start (t0); biotic control of DOR (BC); abiotic control of DOR added with corn maize stalks (AC-CMS) and the same waste incubated with either CMS-immobilized *P. tigrinus* or *Phlebia* sp. (*Pt*-CMS and *P*sp-CMS, respectively); abiotic control treated with polyurethane sponge (AC-PS) and the same waste incubated with PS-immobilized *P. tigrinus* and *Phlebia* sp. (*Pt*-CMS and *Phlebia* sp. (*Pt*-PS and *Ps*p-PS, respectively). Values are the means of three independent experiments, and error bars indicate standard deviations. Statistical pairwise multiple comparisons of data were carried out by the Tukey test: for each plant species, mean values with the same letter are not significantly different ($P \le 0.05$).

fragments were significantly affected by the fungal treatment, as exemplified by the pyrogram of DOR colonized by CMSimmobilized *P. tigrinus* (Figure 3B).

Regardless of the carrier, the abiotic controls of DOR exhibited similar percentages of S, G, and P subunits, in the ranges 3.1-3.2, 9.9-10.7, and 6.3-6.7%, respectively (Table 5). In the same controls, S/G, $(Ac/Ad)_G$, and $(Ac/Ad)_S$ did not significantly differ from one another. Irrespective of the fungus, the relative percent abundances of lignin subunits were significantly modified: most relevant changes were observed for G and S units, the abundances of which were markedly increased. However, S/G ratios in fungal microcosms did not significantly differ from those of the relative abiotic controls with the only exception of CMsupported *Phlebia* sp. $(0.54 \pm 0.02 \text{ vs } 0.30 \pm 0.03, \text{ respectively})$ (Table 5). The $(Ac/Ad)_{S}$ ratios of abiotic controls were generally lower than those of the waste that underwent fungal treatment irrespective of both the fungus and the inoculum carrier; similar results were observed for the (Ac/Ad)_G ratio with the exception of the waste incubated with PS-supported P. tigrinus (Table 5).

PCA performed to analyze TMAH-Py-GC MS showed that around 77% of variability was explained by the first two principal



Figure 2. Correlation between the amount of total phenols added with the waste and *Lepidium sativum* (**A**) and *Lactuca sativa* (**B**) seed germination. To each data point, an Arabic number letter has been assigned to express the related treatment: (1) DOR at start; (2) abiotic control with CMS; (3) abiotic control with PS; (4) biotic control; (5) DOR incubated with free mycelium of *P. tigrinus*; (6) DOR incubated with free mycelium of *P. tigrinus*; (8) DOR incubated with CMS-immobilized mycelium of *P. tigrinus*; (9) DOR incubated with PS-immobilized mycelium of *P. tigrinus*; (10) DOR incubated with PS-immobilized mycelium of *P. tigrinus*; (10) DOR incubated with PS-immobilized mycelium of *P. tigrinus*; (11) distilled water control. Data are expressed in grams of phenols added to 100 g of growth substrate.

components (53.7 and 23.3%, respectively) (Figure 4). Hotelling of scores showed the absence of outliers and, in addition, the observations were clearly separated in different quadrants according to the treatment typology (Figure 4A). In particular, the waste at start and the two abiotic controls were located in the upper and lower right quadrants, respectively. Computation of the contribution of the variables in the loadings plot (Figure 4B) showed that the G%, S% and the S/G ratio had the greatest impact on variability along the first component and P% and $G_n\%$ along the second one (data not shown).

With the only exception of PS-immobilized *Phlebia* sp., the most relevant impact of fungi on DOR carbon pools was a dramatic increase in the amount of TEC (**Table 6**). Fractionation and subsequent purification and quantitation of TEC fractions showed low relative abundance of humic substances (C_{HA+FA}). C_{HA} was significantly higher than the respective abiotic control only in PS-immobilized *P. tigrinus* cultures (126 vs 113 mg g⁻¹, respectively). Despite the low representativeness of C_{HA+FA} in TEC, their absolute amounts were generally higher in fungal cultures than in abiotic controls of DOR except for PS-immobilized *Phlebia* sp. (**Table 6**). To obtain a synoptic and overall indication of the fungal impact on the evolution of organic



Figure 3. Pyrograms of DOR abiotic control added with corn maize stalks (A) and after 4 weeks of treatment at 28 °C with CMS-immobilized *P. tigrinus* (B). For details relative to peak labels, please see Materials and Methods. "LC" indicates those compounds that might derive from both lignin and cellulose (18).

Table 5. Relative Percent Abundances of Syringyl (S), Guaiacyl (G), and *p*-Hydrophenyl (P) Units and Syringyl/Guaiacyl (S/G) and Acid/Aldehyde Ratios [(Ad/Al)_G and (Ad/Al)_S] in Bulk DOR Colonized for 4 Weeks at 28 °C by *Panus tigrinus* or *Phlebia* sp. Supported either on Chopped Maize Stalks (CMS) or on Polyurethane Sponge (PS) and in Its Relative Abiotic Controls

	DOR at start	abiotic controls with		Panus tigrinus supported on		Phlebia sp. supported on	
parameter ^a		CMS	PS	СМ	PS	СМ	PS
G% ^{<i>b</i>}	15.3 AB	9.9 A	10.7 A	15.0 AB	22.3 B	19.1 B	16.9 B
P% ^b	17.0 B	6.3 A	6.7 A	5.9 A	8.1 A	10.2 A	7.4 A
S% ^b	2.9 A	3.1 A	3.2 A	6.0 B	9.1 C	10.3 C	6.8 B
G _n % ^c	43.5 A	51.2 B	52.0 B	55.5 B	56.6 B	48.4 AB	54.5 B
P _n % ^c	48.2 C	32.9 B	32.6 B	21.9 A	20.3 A	25.6 A	23.7 A
S _n % ^c	8.3 A	15.8 B	15.4 B	22.6 C	23.1 CD	26.0 D	21.9 C
S/G	0.20 A	0.31 B	0.30 B	0.41 C	0.41 C	0.54 D	0.40 C
(Ac/Ad) _G	3.95 B	2.39 A	2.48 A	3.36 AB	2.86 A	3.46 AB	5.61 C
(Ac/Ad) _S	0.32 A	0.95 AB	1.35 AB	2.13 B	1.79 B	2.3 B	5.60 C

^a Data are the mean of triplicate runs and row means followed by the same upper case letter did not significantly differ (*P* ≤ 0.05). ^b Data calculated with respect to the total peak areas. ^c Data calculated with respect to the sum of peak areas of P, G, and S subunits.

matter, widely accepted descriptors of the humification process, relating humic substances to different C pools, were calculated. **Table 6** shows that PS-immobilized *Phlebia* sp. was the sole condition where the value of DH, which relates C_{HA+FA} to TEC, was significantly higher than that of the respective abiotic control. By contrast, both HR and HI, relating, respectively, C_{HA} and C_{HA+FA} to TOC, were higher in fungal cultures than in abiotic counterparts (**Table 6**).

DISCUSSION

White-rot fungi are very attractive for decontamination and detoxification purposes (6, 7); their large-scale application,

however, is hampered by several technical constraints, including inocula formulation and mode of application (19). High amounts of inocula (10-20%, w/w), in fact, were needed when free mycelia were incorporated into solid matrices, probably due to friction during mixing (19). As a consequence, different types of carriers to be used in mycoaugmentation have been suggested (20-22).

Although the recalcitrance of DOR is being increasingly identified as one of the major factors hindering microbial approaches aimed at its upgrading (1, 6, 7), the technical aspects associated with fungal inoculum formulation have not yet been given the right consideration. On the other hand, both bioconversion and decontamination of a given solid matrix depend on the

survival and activity of inoculated fungi (21). Spores or mycelial fragments have been reported to be more sensitive to growth inhibition by phenols, such as pentachlorophenol (PCP), than immobilized mycelia (21). The susceptibility of fungi to concentration-dependent growth inhibition by phenols has long been



Figure 4. Principal component analysis of TMAH-py-GC MS data showing scores (**A**) and loadings (**B**) plots. Percent variability explained by each principal component (PC) is shown in parentheses after each axis legend. Abbreviations: DOR, waste at start; AC_CMS abiotic control with control with control with control with polyurethane sponge; *Pt_CMS* and *Psp_CMS*, *P. tigrinus* and *Phebia* sp. immobilized onto corn maize stalks, respectively; *Pt_PS* and *Psp_PS*, *P. tigrinus* and *Phebia* sp. immobilized onto polyurethane sponge. Please see **Table 5** for abbreviations used in plot **B**.

known (23) and should not be neglected when phenol-rich materials such as DOR are considered. In this respect, the different growth responses observed in the present study with free and immobilized inocula might, at least partially, be due to the carrier protective action as also observed by Leštan et al. (21) and Ford et al. (22). In addition, natural immobilization supports, such as lignocellulosic materials, providing nutrients to the mycelial coat might give the fungi an initial competitive advantage over free inocula, leading to increased survival and growth (21, 22). This might explain why in the present study the fungal metabolic activity was highest with CMS-immobilized inocula, whereas with the inert PS, FMIs of immobilized fungi did not differ from those of free mycelia. Use of lignocellulosic substrates coated with alginate-entrapped mycelia conferred the inoculated fungi high antagonistic potential toward indigenous microflora and resistance to growth-inhibiting contaminants likely due to the availability of nutrients arising from the underlying support (20, 21).

In the present study, the use of immobilized inocula led to increased phenols removal with respect to free inocula, regardless of both the fungus and the carrier. However, with regard to DE, a parameter whereby the amounts of phenols removed are normalized to the same amount of biomass, only *P. tigrinus* increased its efficiency with immobilization.

Another interesting aspect associated with the physiology of a phenol-degrading organism is degradation selectivity. In this respect, *Phlebia* sp. exhibited higher DS than *P. tigrinus* with the sole exception of the CMS-immobilized formulate. It is interesting to note that the highest DS values were obtained with immobilized inocula where activity levels of enzymes putatively involved in the oxidation of phenols were significantly higher than the respective free inocula (**Tables 2** and **4**). In this regard, the expression of both laccase and MnP has been reported to be affected by the composition of lignocellulosic mixtures employed in inocula formulation (*22*). Moreover, laccase has been shown to be actively involved in phenols removal from DOR (*24*).

Phytotoxic effects of olive-mill wastes have been suggested to be due to high salt content, relatively low pH, and the presence of other organic components, such as fatty acids (5); phenols, however, are kept as primary determinants of the phytotoxicity of olive residues (1, 2, 6, 11).

The specific mechanisms underlying the phytotoxic action of phenols have not been fully elucidated. However, their toxicity has been attributed to a narcosis mode of action, mainly ascribable to their non-covalent interaction at the membrane level (25). In the present study, immobilized inocula were more effective than free mycelia in reducing phytotoxicity of DOR toward *Lepidium sativum*. On an overall basis, however, *Phlebia* sp. was more effective than *P. tigrinus*; in fact, the latter did not alleviate the DOR phytoxicity toward *Lactuca sativa*. This might be due to the markedly higher efficiency of *Phlebia* sp. in reducing

Table 6. Total Organic Carbon (TOC), Total Extractable Carbon (TEC), Humic and Fulvic Acids Carbon (C_{HA} and C_{FA}, Respectively), Degree of Humification (DH), Humification Ratio (HR), and Humification Index (HI) in DOR at Start (*t*₀) and in the Same Waste That Had Been Incubated with *P. tigrinus* and *Phlebia* sp. Previously Immobilized onto Chopped Maize Stalks or Polyurethane Sponge (*Pt*-CMS and *Ps*p-CMS, Respectively, and *Pt*-PS and *Ps*p-PS, Respectively) and in the Relative Abiotic Controls (AC-CMS and AC-PS)

sample	TOC ^a (mg/g)	TEC ^a (mg/g)	C _{HA} ^a (mg/g)	C _{FA} ^a (mg/g)	$C_{HA+FA}{}^a$ (mg/g)	DH ^a (%)	HR ^a (%)	HI ^a (%)
DOR t_0	495 C	203 A	99.3 A	29.3 A	128.5 A	63.3 C	26.0 A	20.1 A
AC-CMS	479 B	214 A	101.5 A	29.1 A	130.7 A	61.1 C	27.3 A	21.2 A
Pt-CMS	467 A	439 C	110.4 A	46.6 C	157.1 B	35.8 A	33.6 C	23.6 B
Psp-CMS	452 A	420 C	107.2 A	48.3 C	155.4 B	37.0 A	34.4 C	23.7 B
AC-PS	487 B	263 B	112.8 A	27.9 A	140.5 A	53.4 B	28.7 B	23.2 B
Pt-PS	466 A	441 C	125.7 B	24.7 A	150.6 B	34.1 A	32.3 C	27.0 C
<i>P</i> sp-PS	479 B	226 A	98.8 A	37.4 B	136.0 A	60.2 C	28.4 B	20.6 A

^a Data are the mean of three independent experiments: column means followed by the same upper case letter are not significantly different (P ≤ 0.05).

Article

ethyl acetate-extractable phenols (Tables 2 and 3). This fraction mainly encompasses the waste's monomeric phenols, which, by virtue of their reduced molecular weight and lipophilicity, have better access to the plant membrane than the polymeric ones (11). This hypothesis is supported by a recent study reporting that Lepidium sativum germinability was not negatively affected by the residual aqueous fraction of DOR obtained through extensive ethyl acetate extraction (2). Moreover, a clear relationship between trans-cinnamic derivatives, such as p-coumaric acid, and abscissic acid (ABA) on the inhibition of Lactuca sativa germination has been reported (26). With this specific regard, P. tigrinus was significantly less effective than Phlebia sp. in the removal of p-coumaric acid from DOR. In addition, P. tigrinus actively produced ABA in olive-mill waste-based medium (27). The biological conversion of lignocellulosic residues by white-rot fungi leads to the formation of humic-like substances, the amount of which depends on the balance between polymerization and depolymerization reactions (28) and on the variable degradation ability of these organisms toward humic acids (29). In the present study, an overall increase in the amount of C_{HA+FA} was found in DOR that had been incubated with immobilized fungi with respect to abiotic controls; this effect, however, was associated with a concomitant drastic increase in the amount of TEC leading to lower DH values, thus indicating that the humification process in DOR was just at its early stage. This was also confirmed by the higher abundance of CFA than CHA and by the low percent abundance of the latter in TEC. This indication was supported by TMAH-Py-GC MS analyses of the waste, which showed that the S/G ratio did not change upon fungal treatments, although it was expected to be decreased in the presence of an extensive delignification (12, 13). However, the generalized increase in the $(Ac/Ad)_{S}$ and, in some cases, the $(Ac/Ad)_{G}$ ratio in DOR incubated with fungi suggested the occurrence of oxidative cleavage of the C_{α} — C_{β} bonds in the propenyl side chain of lignin subunits; this is one of the main mechanisms involved in lignin depolymerization (13). Interestingly, the waste exhibited a significant presence of H monolignols and very low S/G ratio. The high abundance of P subunits in DOR is interesting and anomalous, because it arises from Olea europea L., a member of dicotyledonous angiosperms, the ligning of which are mainly composed by G and S moieties (30). The S/G ratio has been suggested to be an important determinant for the susceptibility of a lignocellulosic material to biodegradation, the extent of which increases as the ratio increases (13). The rationale is that the S units, besides being characterized by redox potentials lower than the G units, are connected via relatively labile ether bonds in lignin. Conversely, G monolignols tend to form highly condensed lignin substructures with a predominance of both stable biphenyl and other carbon-carbon linkages, which are less susceptible to biodegradation (30). Therefore, the widely reported recalcitrance of the waste to biological degradation (1) might be explained both by these structural characteristics and by the large abundance of endocarp fragments, where the majority of lignin is located, with low surface area exposed to microbial attack.

ABBREVIATIONS USED

 $(Ad/Al)_G$, guaiacyl acid/aldehyde ratio; $(Ad/Al)_S$, syringyl acid/aldehyde ratio; Ar/Al, aromatic/aliphatic ratio; AWSP, acetone/water-soluble phenols; C_{FA}, fulvic acid carbon; C_{HA}, humic acid carbon; CMS, chopped maize stalks; DE, depheno-lization efficiency; DH, degree of humification; DOR, dry olive-mill residue; DS, dephenolization selectivity; EAEP, ethyl acet-ate-extractable phenols; FA, fatty acids; FMI, fungal metabolic index; G, guaiacyl subunits; P, *p*-hydroxyphenyl subunits; HI,

humification index; HR, humification ratio; IC₅₀, concentration required for 50% germinability inhibition; MnP, manganese peroxidase; OM, organic matter; PCA, principal component analysis; PS, polyurethane sponge; RP-HPLC, reversed-phase high-performance liquid chromatography; S, syringyl subunits; S/G, syringyl/guaiacyl ratio; TEC, total extractable carbon; TOC, total organic carbon; TMAH, tetramethylammonium hydroxide; TPEP, two-phase extraction process; WRF, white-rot fungi.

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