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Biodegradation of oxytetracycline by *Pleurotus ostreatus* mycelium: a mycoremediation technique

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1. Introduction

Oxytetracycline (OTC) is an antimicrobial drug widely used in intensive farming to treat enteric and respiratory diseases [1,2]. Due to oral administration and its pharmacokinetic, OTC dosage is in the range of $g kg^{-1}$ medicated feeds and it is found in animal waste at concentrations of $mg kg^{-1}$ [3]. As a consequence of waste disposal, the drug enters into the environmental compartments (water, sediments, soil) and can contaminate trophic nets [4–6], determining carry over in food and negative effects on natural systems [7,8], including effects on the microbial community structure and selection of antibiotic-resistant strains [9–11].

To avoid the negative effects, drug degradation and/or removal are a possible solution that can be obtained by bioremediation techniques. Fungi have not extensively investigated for biotransformation of antibiotics [12]. Some of them have been applied in polycyclic aromatic compounds bioremediation and dyes decolorization [13–15]. Wen et al. [16,17] studied the degradation of tetracycline and OTC by crude ligninolytic enzymes (lignin peroxidase and manganese peroxidase) prepared from *Phanerochaete chrysosporium*, with positive results. Their studies were conducted *in vitro* and did not include tests on another important ligninolytic enzyme, the laccase, aplenty produced by white-rot fungi.

ABSTRACT

Oxytetracycline (OTC) is administered in high doses to livestocks and enters the environmental compartments as a consequence of animal waste disposal. As a first step in setting up a useful mycoremediation technique, an OTC lab degradation test was performed in liquid medium using the ligninolytic fungus *Pleurotus ostreatus*. OTC disappearance in culture medium was clearly evident as early as the third day of exposure onwards, with an almost complete removal after 14 d. The drug removal was mediated by fungal absorption in the mycelia, where the OTC molecule underwent a degradation step, as demonstrated by mass spectrometry analyses. A putative degradation product, ADOTC (2-acetyl-2-decarboxamidooxytetracycline) is proposed. Experimental conditions excluded OTC abiotic degradation; the degradation by extracellular laccase was also experimentally discarded.

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In a previous study, Galli et al. [18] utilized *P. ostreatus* SMR684 in mycoremediation experiments for the degradation of creosote components. This strain is known to produce laccase both in the presence of lignocellulosic substrates and constitutively, in some phases of its life cycle [19].The aim of this study is to investigate the ability of the fungus *P. ostreatus* SMR684 to degrade OTC. To this end in a laboratory model, living mycelia were challenged with different OTC concentrations. A positive result could represent the first step of a mycoremediation technique on a pilot scale.

2. Materials and methods

2.1. Organism and culture conditions

P. ostreatus (Jacq.) P. Kumm. mycelium, strain SMR684, is stored in minimal growth at +5 °C in the Edible Saprophytic Mushrooms collection of IBAF-CNR, Rome, Italy. A small fraction of mycelium was grown for 2 d in a 250 mL Erlenmeyer flask containing 100 mL of a sterilized 3% malt extract liquid culture medium (Liofilchem s.r.l., Roseto, TE, Italy) on a rotary shaker (100 rpm), under controlled laboratory conditions at $T=25 \pm 1$ °C and Relative Humidity (RH)=70 ± 5%.

Pre-inoculum culture was started by aseptically homogenizing the mycelium in an omni-mixer for 3 s and transferring 50 mL of mycelial suspension into a flask containing 500 mL of 3% malt extract. The mycelium grew in the controlled laboratory conditions for 3 d then aliquots of this culture were used as inocula.



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2.2. The experimental plan

The experimental plan was organized on 6 batches: (1) culture medium (3% malt extract); (2) culture medium + 50 μ g mL⁻¹ OTC; (3) culture medium + 100 μ g mL⁻¹ OTC; (4) culture medium + *P. ostreatus*; (5) culture medium + *P. ostreatus* + 50 μ g mL⁻¹ OTC; (6) culture medium + *P. ostreatus* + 100 μ g mL⁻¹ OTC. Batches from #1 to 4 are controls, #5 and #6 are test batches. Each batch was constituted by 18 Erlenmeyer flasks (3 replicates for each day of sampling). The entire experiment has been repeated three times.

A sterilized solution $(0.2 \,\mu m \text{ filter})$ of OTC (Sigma, Milan) in deionised water was added to the test cultures up to the final concentrations in the culture media of 50 and $100 \,\mu g \, m L^{-1}$. Aliquots of 6 mL of the pre-inoculum culture (corresponding to 14.4 mg of mycelium, dry weight, d.w.) were used as inocula.

All flasks were incubated at 25 ± 1 °C and $70 \pm 5\%$ RH. Samples of each batch were collected at the beginning of the trial (T0) and after 2 (T2), 3 (T3), 7 (T7), 10 (T10) and 14 d (T14). To avoid OTC photodegradation, flasks were entirely covered by aluminium sheets and all experimental procedures were performed under dark. Different samples were obtained: from #1 to 3 batches only culture media; from #4 to 6 both mycelia and culture media. Mycelia were separated from culture medium by gauze filtration. Culture media were collected and immediately stored at -20 °C until chemical and enzymatic analyses. Then mycelia were rinsed with distilled water, freeze dried, weighed and stored at room temperature until chemical analysis.

2.3. Enzymatic activity

2.3.1. Laccase in the culture medium

Culture medium aliquots of 2.5 mL from flasks of batch #4 and 6 were centrifugated at 9000g for 20 min and 4 °C and the supernatant was used for the laccase assay. It was performed at 25 °C in 2.9 mL of 100 mM sodium acetate buffer, pH 4.5, containing 2 mM 2,6-dimethoxyphenol, plus 0.1 mL of the culture medium supernatant (3 mL final volume). The oxidation of 2,6-dimethoxyphenol was measured by spectrophotometry at 468 nm (ε_{468} = 49600 M⁻¹ cm⁻¹; Perkin-Elmer Lambda 3B spectrophotometer). One unit of laccase activity was defined as the amount of enzyme required to oxidise 1 µmol substrate min⁻¹ [20]. Results were analyzed by One-Way ANOVA.

2.3.2. Effect of purified laccase on OTC degradation

Direct OTC degradation by purified commercial laccase (from *Trametes versicolor*, 21.8 U mg⁻¹, Sigma, Milan, Italy) was evaluated by incubating 50 or 100 μ g mL⁻¹ OTC with laccase (0.1 U mL⁻¹ in 100 mM sodium acetate buffer, pH 4.5). Reaction was carried out under dark at 25 °C in a final volume of 20 mL in amber flask, under continuous stirring. Control test was performed on OTC and buffer. Degradation was evaluated periodically up to 96 h. OTC quantification was performed by direct injection of each sample in a LC–UV/DAD system (see Section 2.4.3).

2.4. Chemical analysis

2.4.1. Extraction

Mycelia were ground in a mortar, 100 mg d. w. were poured into a 50 mL Falcon tube added with 100 μ L of a methanol solution of Doxycycline hyclate (DXC, Sigma, Milan, Italy) 100 μ g mL⁻¹ and extracted twice with 10 mL of 0.05 M sodium acetate. After homogenization, sonication and mechanical agitation (15 min), the sample was centrifuged (5000 g, 5 min, room temperature) and the pooled supernatants were cleaned-up on Metal Chelate Affinity Chromatography (MCAC) columns. Culture medium from each sample was filtered through a 0.45 μ m sterile membrane; 1 mL



Fig. 1. Growth of *P. ostreatus* mycelium in control batch (green), $50 \ \mu g \ mL^{-1}$ OTC (blue) and $100 \ \mu g \ mL^{-1}$ OTC (red) exposed batches. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of the article.)

of filtered sample was added with 100 μ L DXC methanol solution (100 μ g mL⁻¹) and cleaned-up on MCAC columns.

2.4.2. Purification

MCAC columns were prepared by filling 15 mL chromatography glass columns with 1.5 mL of Chelating Sepharose Fast Flow (GE Healthcare, Uppsala, Sweden). The sorbent was washed with 2 mL water: ethanol (80:20) solution and 6 mL water, conditioned with 3 mL of 0.01 M CuSO₄ solution, then washed twice with 2 mL water. Columns were loaded with samples and then washed with 2 mL of water, then 2 mL methanol and 2 mL water. Elution was achieved with the application of 3 mL McIlvaine buffer (pH 4.5). 20 μ L eluate were injected into the HPLC–UV/DAD system.

In LC–MS analysis, 3 mL samples (in McIlvaine buffer) were concentrated with SPE (Solid-Phase Extraction) polymeric columns (Oasis HLB, 60 mg 3 mL, Waters), preconditioned with 3 mL methanol and 3 mL water. After sample loading the column was washed with 3 mL of water, then vacuum dried. Elution was achieved with the application of 3 mL methanol, sample were dried under nitrogen flow and reconstituted in 100 μ L methanol. 20 μ L sample were injected onto the LC–ESI-MS system.

2.4.3. HPLC–UV/DAD analysis

An Agilent 1100 series HPLC system equipped with a RP XTerra MS C₈ column (5 μ m, 2.1 mm \times 150 mm) and photodiode array detector (set at 365 nm) was used. Mobile phases were: A, 1% acetic acid in water (v/v) and B, acetonitrile HPLC grade. Flow rate: 0.3 mL min⁻¹. Course gradient was: 0'-2.5' 2% B, 2.5'-10.0' 100% B, 10.0'-12.0' 100% B, 12.0'-13.00' 2% B, 13.0'-15.00' 2% B. The estimated Limit of Detection (LOD) was 1 ppm.

2.4.4. LC-ESI-MS analysis

Liquid chromatography–electrospray ionization-mass spectrometry (LC–ESI-MS) analyses were performed with a Agilent 1100 series LC-MSD Trap mass spectrometer coupled to the HPLC system previously described, working at the same conditions. Analytes were detected using ESI in positive mode.

3. Results

3.1. Fungal growth

The growth of each experimental culture at different times is reported in Fig. 1 as mycelia dry weight. Both control and test





Fig. 2. LC–UV/DAD analysis of OTC in mycelia from: control culture at T3 (first row); cultures exposed to OTC 50 μg mL⁻¹ at T3 (second row, left) and T14 (third row, left); cultures exposed to OTC 100 μg mL⁻¹ at T3 (second row, right) and T14 (third row, right). OTC = Oxytetracycline; DXC = Doxycycline (internal standard).

cultures showed active growth during the entire experimental period. After 14 d (T14) control cultures showed a 50-fold increase, from the initial 0.14 mg mL⁻¹ of the inoculum to a final mean weight of 7.06 mg mL⁻¹. The 50 μ g mL⁻¹ OTC culture reached the same final weight increase at T14, although weight is clearly lower than control until T10. Over the entire test the 100 μ g mL⁻¹ OTC culture showed lower fungal weight than the control. At T14 fungal growth reached a 36-fold increase from the initial inoculum, to a final mean weight of 5.13 mg mL⁻¹.

3.2. OTC degradation

OTC in the different batches and at different times was quantified by LC–UV/DAD analysis. The chemical analysis of the extract from control culture media and mycelia (batches #1 and 4) excluded the presence of interfering peaks in the area of interest of the chromatograms. Analysis of culture media containing 50 or 100 μ g mL⁻¹ OTC without mycelium showed almost the same OTC concentrations from T0 to T14, demonstrating that OTC abiotic degradation did not occur in neither batch #2 or 3 under these experimental conditions. On the contrary, the analysis of culture media and mycelia extracts from OTC exposed cultures (batches #5 and 6) showed a dramatic reduction of OTC concentration in the course of the experiment (Fig. 2 and Fig. 3).

In particular, culture media of batch #5 showed a dramatic OTC reduction in the first 7 d and a further reduction in the last 7 d, up to an almost complete absence of OTC residues at T14 ($0.6 \ \mu g \ mL^{-1}$). Contemporary, in batch #5 mycelia, T2 and T3 extracts showed an increase of OTC content, followed by a progressive decrease up to the complete absence at T14 (Fig. 3).



Fig. 3. OTC concentrations determinated by HPLC–UV/DAD in culture medium or mycelium at different times in 50 and $100 \,\mu g \, mL^{-1}$ OTC exposed batches.

A comparable trend of OTC degradation was found also in batch #6. In the culture media, a sharp reduction of the initial concentration in the first 7 d and a further limited reduction in the last 7 d occurred; as a result a very low OTC concentration was found at T14 ($2.8 \ \mu g \ m L^{-1}$). In mycelia extracts an initial increase of OTC content was found (T2–T3), followed by a progressive significant decrease that reduced the OTC residue at T14–7.6 $\mu g \ per 100 \ mg \ mycelium$ (d.w.; Fig. 3).

3.3. Laccase activity in the culture medium

The laccase activity during the degradation process was measured in the culture media of control (batch #4) and OTC exposed culture (batches #6; Fig. 4). *P. ostreatus* showed the production of extracellular laccase in the course of the trial, the maximum values being reached at T4–T5. No significant differences were obtained between control and OTC 100 μ g mL⁻¹, although a slightly higher activity was recorded from T3 to T5 in the presence of OTC.

3.4. Effect of purified laccase on OTC degradation

The degradation capability of the extracellular enzyme alone on OTC was tested by an *in vitro* test, coupled to OTC quantification. OTC solutions prepared according to the experimental conditions



Fig. 4. Laccase activity measured in the culture medium of control and 100 $\mu g\,mL^{-1}$ OTC treated batch.

Table 1

Percentage of OTC degradation due to purified commercial laccase. The results were produced by comparing the amount of OTC degraded in the batches with or without added laccase at each time.

Time (h)	OTC 50 $\mu g m L^{-1}$ (%)	OTC 100 µg mL ⁻¹ (%)
0	2.2	1.0
1	1.8	0.5
2	2.4	0.8
3	2.3	1.3
5	3.0	0.4
24	2.4	0
48	4.5	0.9
72	6.6	0
96	5.0	1.6

reported in Section 2.3.2 were incubated with or without laccase. No significant differences in OTC degradation were measured in the two batches (Table 1).

3.5. Products of OTC fungal degradation

OTC degradation has been followed within the mycelial matrix using LC–ESI-MS full scan analysis of the extracts to detect the possible presence of degradation products. The results show three partially overlapping peaks (Fig. 5, coloured chromatogram). The first one (red) shows an OTC precursor ion $[M+H]^+ m/z$ 461, the second one (green) shows a precursor ion $[M+H]^+ m/z$ 460, that



Fig. 5. Chromatogram of the extracted precursor ions [M+H]⁺ m/z 461(in red, OTC), 460 (in green) and 445 (in black, DXC) obtained by LC–ESI-MS. The inset shows the full scan spectrum in the area where the three chromatographic peaks are overlapped. (For interpretation of the references to color in this figure legend, the reader is referred to the web version of the article.)



Fig. 6. Chemical structure of OTC (a) and ADOTC, 2-Acetyl-2-Decarboxamido-Oxytetracycline (b).

could be a degradation product of OTC, and the third one (black) a DXC precursor ion $[M+H]^+ m/z$ 445 (internal standard) (Fig. 5, inset).

4. Discussion

The proposed experimental model allows to demonstrate that (i) *P. ostreatus* is able to survive and grow in the presence of OTC, and (ii) the growing fungus actively removes (and degrades) the drug from the culture medium.

The growth dynamics of control and OTC exposed cultures shows that *P. ostreatus* mycelia survive and grow even in the presence of considerable amount of drug, although the growth rate is reduced at the higher drug concentration. In fact, at $50 \,\mu g \, m L^{-1}$, the rapid removal/degradation of the drug allows the mycelium to grow reaching the same final values of the control. At $100 \,\mu g \, m L^{-1}$, the mycelium starts a fast growth just after the removal/degradation of a considerable amount of drug.

While pH and UV can induce epimerization and isomerization reactions in the molecule [21], at our experimental conditions OTC does not undergo abiotic modifications. Drug removal (and degradation) from the culture medium is carried out by the growing fungus. In fact, after 7 d OTC in the culture media drops to 23% of the spiked drug, at both concentrations and the reduction still continues reaching very low amounts (<3%) in the subsequent 7 d of experiment. While OTC is removed by the culture media, appreciable amounts of drug can be found within the mycelium (after 3 d exposure), demonstrating the OTC uptake by the fungus. Thus, the OTC reduction in the culture media has to be ascribed to the fungus, able to uptake the drug. The uptake is soon followed by an efficient degradation of the drug, being the amounts of OTC detected in the mycelia a spare fraction of the drug supplied.

To date these results on OTC cannot be compared with other experimental data, although *P. ostreatus* is known to absorb and degrade recalcitrant compounds [18,22]. In a study devoted to fluoranthene degradation, as a first step the compound was completely absorbed by the *P. ostreatus* mycelium (15 d) and then degraded to 50% of the spiked amount in 40 d [23]. In a study on phenanthrene degradation using *P. ostreatus*, phase I and II metabolites were screened; the results indicated that the phase II enzymes, as Glutatione S-transferase, were not important for the elimination of hydroxylated phenanthrene [24].

As regard the involvement of laccase in OTC degradation under our experimental conditions, degradation test showed that laccase by itself cannot be considered as the prior responsible for fungal degradation of OTC, being unable to degrade OTC in the absence of mycelia. This result is in agreement with the experiments of Wen et al. [16,17], which demonstrated the degradation activity of a crude extract in which several enzymes and redox mediators were present.

The drug degradation process within the mycelial matrix has been shown in its early phases: LC–ESI-MS analyses allowed to follow the first degradation step of mycelia on the OTC molecule (Fig. 6a). In fact, in addition to the expected OTC precursor ion $([M+H]^+ m/z 461)$, the mass spectrum revealed the presence of another precursor ion $([M+H]^+ m/z 460)$; this additional precursor ion could be interpreted as 2-acetyl-2-decarboxamido-oxytetracycline (ADOTC; MW=459; Fig. 6b). This compound is known to be a side product of OTC production by *Streptomyces rimosus* [25]. Its presence was analytically excluded in the standard solutions as impurity, but was revealed in the treated batches, only in the first days of the experiment. ADOTC has lower antibacterial potency on environmental relevant bacteria and higher lipophilicity than OTC [25]. These chemical characteristics determine a lower toxicity and environmental mobility, fulfilling the aim of our bioremediation approach.

5. Conclusions

This approach represents a first and successful step towards a feasibile and unexpensive bioremediation technique to avoid OTC spreading into environmental matrices.

P. ostreatus SMR684 represents a useful tool for OTC mycoremediation from contaminated matrices: the fungus is not only able to survive and grow in the presence of significant amount of drug, but also almost completely degrades the drug in few days. The drug is absorbed by the mycelia, where the OTC molecule undergoes a degradation step, as demonstrated by mass spectrometry analyses. In the treated mycelia no OTC-like residues are detected at the end of the experiment.

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