

## Biotransformation of Malachite Green by the Fungus *Cunninghamella elegans*

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**The filamentous fungus *Cunninghamella elegans* ATCC 36112 metabolized the triphenylmethane dye malachite green with a first-order rate constant of 0.029  $\mu\text{mol h}^{-1}$  (mg of cells)<sup>-1</sup>. Malachite green was enzymatically reduced to leucomalachite green and also converted to N-demethylated and N-oxidized metabolites, including primary and secondary arylamines. Inhibition studies suggested that the cytochrome P450 system mediated both the reduction and the N-demethylation reactions.**

Malachite green, an N-methylated diamino-triphenylmethane dye, has been widely used as the most efficacious antifungal agent in the fish farming industry (26). It is also used extensively in textile industries for dyeing nylon, wool, silk, leather, and cotton (10). Although malachite green is not approved by the U.S. Food and Drug Administration, its worldwide use in aquaculture will probably continue due to its relatively low cost, ready availability, and efficacy (26); therefore, potential human exposure to malachite green could result from the consumption of treated fish (2) and from working in the dye and aquaculture industries. Malachite green is highly toxic to mammalian cells; it promotes hepatic tumor formation in rodents and also causes reproductive abnormalities in rabbits and fish (13, 24). The structural similarity of malachite green to other carcinogenic triphenylmethane dyes also raises suspicion of carcinogenicity; gentian violet (crystal violet) is a thyroid and liver carcinogen in rodents (17), and pararosaniline is a bladder carcinogen in humans (7). Based on the potential for adverse human health effects, the U.S. Food and Drug Administration nominated malachite green as a priority chemical for carcinogenicity testing by the National Toxicology Program in 1993 (10). These studies are presently being conducted at the National Center for Toxicological Research, Jefferson, Ark.

From an environmental standpoint, there is concern about the fate of malachite green and its reduced form, leucomalachite green, in aquatic and terrestrial ecosystems, since they occur as contaminants (6, 21) and are potential human health hazards. Studies on the biodegradation of triphenylmethane dyes have focused primarily on the decolorization of dyes via reduction reactions (4, 19, 22, 23, 25). Intestinal microflora were shown to reduce crystal violet (18) and malachite green (16) to their respective leuco derivatives. The fungal metabolism of these compounds was first reported by Bumpus and Brock (5). The white rot fungus *Phanerochaete chrysosporium*, grown under ligninolytic conditions, was shown to metabolize crystal violet to three metabolites by sequential N demethylation of the parent compound, which was catalyzed by lignin peroxidase. They also reported (5) that nonligninolytic cultures

of *P. chrysosporium* could also degrade crystal violet, although the N-demethylation products were not found under nonligninolytic conditions, suggesting that another mechanism for degrading crystal violet existed in this fungus. The present study was conducted to determine whether the filamentous fungus *Cunninghamella elegans*, which has been used as a microbial model for mammalian xenobiotic metabolism (1) as well as for the biodegradation of environmentally relevant chemicals (8), had a mechanism in triphenylmethane dye metabolism different from that of *P. chrysosporium*. *C. elegans* is capable of metabolizing a wide range of compounds, especially by N demethylation and N oxidation (14, 20, 27, 28). Little is known about the potential of nonligninolytic fungi to metabolize triphenylmethane dyes. This paper describes the metabolic fate of malachite green by cultures of *C. elegans*.

Biotransformation experiments were performed by the addition of malachite green (97% dye content; Aldrich Chemical Co., Milwaukee, Wis.) or leucomalachite green (Aldrich Chemical Co.) to 48-h-old cultures of *C. elegans*. Culture conditions were as described previously (20). Leucomalachite green was dissolved in dimethylformamide before addition. The data are averages based on three separate experiments performed with duplicates. After 5 days of incubation, fungal mycelia were removed by filtration and extracted with ethyl acetate (five times, each time with 100 ml). The supernatant was also extracted with ethyl acetate. The ethyl acetate extracts were then dried over anhydrous  $\text{MgSO}_4$  and evaporated in vacuo. The dried sample was dissolved in 10 ml of solution containing acetonitrile (60%) and 50 mM ammonium acetate (pH 4.5) (40%) for analysis by high-performance liquid chromatography (HPLC) and HPLC-mass spectrometry (MS).

Reverse-phase HPLC was performed with a Hewlett-Packard (Palo Alto, Calif.) 1050 series component system equipped with a photodiode array detector. Samples were resolved on a Spherisorb S5 nitrile column (4.6 by 250 mm; particle size, 5  $\mu\text{m}$ ) with a  $\text{PbO}_2$  post-column (4.6 by 10 mm) to detect nonchromatic leucomalachite green and its derivatives at 618 nm following oxidation to chromatic forms (3). The metabolites were eluted at a flow rate of 1.0 ml/min with a linear gradient running from 30% to 90% B (solvent A, 50 mM ammonium acetate, pH 4.5; solvent B, acetonitrile) for 30 min. An isocratic solvent system (solvent A/solvent B ratio = 40:60) was also used when the disappearance of malachite green was moni-

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tored. Conditions for liquid chromatography-atmospheric pressure chemical ionization-MS analysis were as described previously (12).

Cultures of *C. elegans* transformed malachite green, up to 54  $\mu\text{M}$ , with a first-order rate constant of  $0.029 \mu\text{mol h}^{-1}$  (mg of cells) $^{-1}$ . Apparently 85% of malachite green in culture flasks (81  $\mu\text{M}$ ) had disappeared after 24 h. A concentration of 108  $\mu\text{M}$  malachite green inhibited fungal growth, and biotransformation did not occur. The absorption spectra of samples removed during biotransformation indicated that the wavelength (618 nm) at which malachite green exhibits its chromatic feature shifted to 608 nm after 8 h of incubation. These results suggested that malachite green might be undergoing N demethylation, since the N-demethylation products have absorption maxima at wavelengths lower than that of malachite green (B. P. Cho, personal communication). The loss of color was observed during incubation, suggesting that malachite green was reduced to its leuco- form (16). To confirm this observation, the metabolites from ethyl acetate extracts of *C. elegans* cultures incubated with malachite green and leucomalachite green were analyzed by HPLC in combination with atmospheric pressure chemical ionization-mass spectrometry. Figure 1 shows reconstructed molecular ion chromatograms from the samples extracted from the fungal cells after 5 days of incubation. Under these conditions, the mass spectra consisted primarily of molecular ions (protonated molecules for leucomalachite green and the demethylated derivatives and cationic molecules for malachite green and its derivatives). Based on previous reports (11, 12), these peaks correspond to malachite green ( $m/z$  329) and its mono-, di-, and tri-desmethyl derivatives ( $m/z$  315, 301, and 287, respectively) and leucomalachite green ( $m/z$  331) and its mono-, di-, tri-, and tetra-desmethyl derivatives ( $m/z$  317, 303, 289, and 275, respectively). The metabolites extracted from the culture supernatants were similar to those obtained from mycelium-extracted samples, except for malachite green N-oxide ( $m/z$  345; retention time, 9.21 min), which was detected only in the mycelia. Control experiments with autoclaved cells did not produce a significant amount of metabolites. Only leuco- derivatives were observed as the final products of biotransformation after a prolonged incubation time (10 days), suggesting that the N-demethylated malachite green metabolites were also reduced to their corresponding leuco- derivatives. When leucomalachite green was used as the initial substrate, identical patterns of metabolites (mono-, di-, tri-, and tetra-desmethyl leucomalachite green) were observed.

The microsomal fraction from *C. elegans*, which was prepared as described previously (9), also appeared to mediate the transformation of malachite green. The incubation mixtures contained the following components in a total volume of 2 ml: 0.1 mg of malachite green, 1 mM NADPH, and 2.5 mg of microsomal protein in 50 mM sodium phosphate buffer, pH 7.0. Desmethyl and di-desmethyl malachite green and leucomalachite green were detected by HPLC. Boiled microsomal protein did not produce any demethylated metabolites. Leucomalachite green and its demethylated metabolites were not formed in the absence of NADPH, although demethylated metabolites of malachite green were still produced.

Cytochrome P450 inhibitors, such as 1-aminobenzotriazole (2 mM), metyrapone (2 mM), and SKF 525-A (1.5 mM), retarded biotransformation of malachite green. Metyrapone

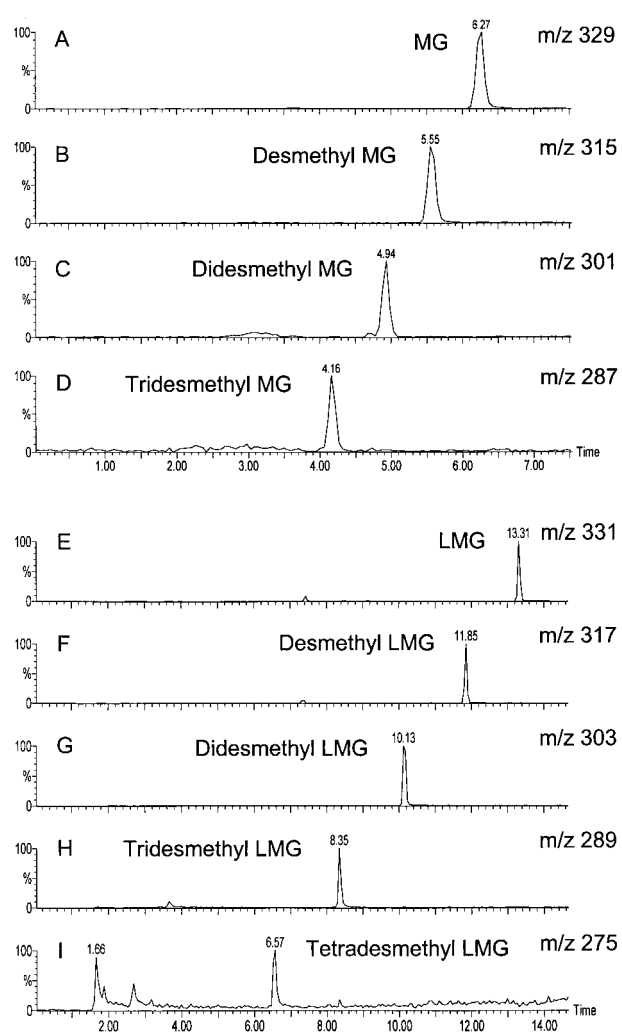


FIG. 1. LC-atmospheric pressure chemical ionization-mass spectrometry molecular ion chromatograms obtained at 20 V from an ethyl acetate extract of *C. elegans* incubated with 32  $\mu\text{M}$  malachite green for 5 days. (A)  $m/z$  329, malachite green (retention time, 6.27 min); (B)  $m/z$  315, desmethyl malachite green (retention time, 5.55 min); (C)  $m/z$  301, didesmethyl malachite green (retention time, 4.94 min); (D)  $m/z$  287, tridesmethyl malachite green (retention time, 4.16 min); (E)  $m/z$  331, leucomalachite green (retention time, 13.31 min); (F)  $m/z$  317, desmethyl leucomalachite green (retention time, 11.85 min); (G)  $m/z$  303, didesmethyl leucomalachite green (retention time, 10.13 min); (H)  $m/z$  289, tridesmethyl leucomalachite green (retention time, 8.35 min); (I)  $m/z$  275, tetradesmethyl leucomalachite green (retention time, 6.57 min)

completely inhibited the reactions; 1-aminobenzotriazole inhibited the reactions by 67%, and SKF 525-A inhibited them by 70%. This suggested that the cytochrome P450 system of *C. elegans* mediated the N-demethylation reaction as well as the reduction of malachite green to leucomalachite green.

Previous studies (5, 22) demonstrated that the white rot fungus *P. chrysosporium* employed extracellular lignin peroxidases under ligninolytic conditions to decolorize crystal violet by sequential N demethylation. However, the present study shows that the nonligninolytic fungus *C. elegans* has multiple pathways to transform triphenylmethane dyes by intracellular cytochrome P450(s) which mediate(s) both the reduction and

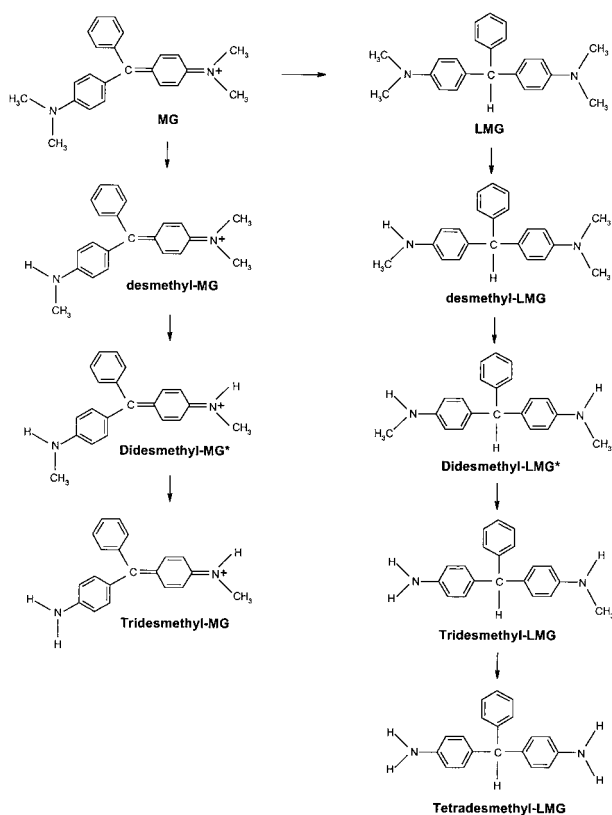


FIG. 2. Proposed mechanism for the metabolism of malachite green (MG) and leucomalachite green (LMG) by *C. elegans*. The asterisk indicates that unsymmetrical didesmethyl MG and LMG are not shown.

the N demethylation (Fig. 2). This study demonstrated that the decolorization of malachite green by *C. elegans* could be attributed mainly to its reduction to leucomalachite green since the demethylated metabolites of malachite green still exhibit absorption at 618 nm. The reduction of crystal violet by rat liver microsomes was shown to be catalyzed by a cytochrome P450 monooxygenase system via a one-electron reaction (15). The present study also suggested that *C. elegans* employs cytochrome P450 for the reduction of malachite green, because the cytochrome P450 inhibitors used in this study, especially metapyrone, clearly inhibited the reduction. Our study also demonstrated that this fungal system produced metabolite profiles similar to those observed in rat liver (11). Thus, *C. elegans* is a suitable microbial model for triphenylmethane dye metabolism and will be used to produce significant quantities of metabolites for toxicological evaluation.

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