

Full Length Research Paper

Biotransformation of coumarins by *Cunninghamella elegans*

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Microbial biotransformation of coumarin (I) was undertaken using a battery of twenty five microorganisms. Among them, the fungus *Cunninghamella elegans* NRRL 1392 was the sole microbe that efficiently metabolized coumarin (I) into umbelliferone (II), 3, 4-dihydrocoumarin (V) and transcinnamic acid (VI). Umbelliferone was also biotransformed into 6, 7-dihydroxycoumarin (aesculetin, VII) and p-coumaric acid (VIII). On the other hand, warfarin (III) was biotransformed into 4'-hydroxywarfarin (IX) and also dicoumarol (IV) was transformed into 4-hydroxycoumarin (X) by the same microorganism. The structures of the metabolites were established using only one property and spectroscopic techniques including melting points, ¹H NMR, ¹³C NMR and mass spectroscopy. The cytotoxic effect of coumarin substrates and their metabolites was also investigated.

Key words: Biotransformation, coumarins, *Cunninghamella elegans*, cytotoxic.

INTRODUCTION

Coumarin ring system (benzo- α -pyrone) (Dean, 1963) is widely encountered in natural products in many plant families, such as Umbelliferae, Rutaceae and others (Murray et al., 1982; Berenbaum, 1991). Coumarin (I) is used extensively as perfume constituent (Trumble et al., 1990), while its derivatives umbelliferone (II), warfarin (III) and dicoumarol (IV) may exhibit activity against several types of animal tumors (Beier and Nigg, 1994). Furthermore, umbelliferone (II) demonstrated an antioxidant activity (Beier and Oerti, 1983), while dicoumarol (IV) and warfarin (III) possessed a potential antineoplastic (Jamal and Casley Smith, 1989) as well as anticoagulant (Musajo and Rodighiero, 1962) effects, respectively. The use of microbial transformation in

simulation of mammalian drug metabolism was firstly reported by Smith and Rosazza (1975) to insight the drug metabolic pathway, mechanism of action and toxicity. These studies were based on the parallels, which exist between phase-1 and phase-2 reactions in both microbial and mammalian systems (Van Scott, 1976; Scott et al., 1976) and had led in overcoming several complications of low yield, high cost and impure metabolites appeared during the studies of drug mammalian metabolism. Recently, this approach is used in the synthesis of new biologically active relevant products (Wackett et al., 1995).

Bioconversion of some coumarins derivatives into new therapeutic products using several microorganisms was

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Table 1. Results of screening coumarin (I) biotransformation with microbial cultures.

Microorganism	Screening results	Incubation period
<i>Aspergillus alliceous</i> NRRL315	-	10
<i>Aspergillus flavipes</i> ATCC 11013	-	10
<i>Aspergillus niger</i> NRRL 599	+	5
<i>Aspergillus niger</i> NRRL 2295	+	5
<i>Aspergillus niger</i> ATCC 10549	+	5
<i>Aspergillus ochraceous</i> NRRL 398	+	7
<i>Aspergillus ochraceous</i> NRRL 405	-	7
<i>Candida albicans</i> (lab isolate)	+	7
<i>Cunninghamella bainari</i> UI 3605	+	7
<i>Cunninghamella blackesleeana</i> 8688a	-	7
<i>Cunninghamella blackesleeana</i> MR 398	-	7
<i>Cunninghamella echinulata</i> NRRL 1382	+	7
<i>Cunninghamella elegans</i> NRRL 1392	+++	4
<i>Gymnascella citrine</i> NRRL 6050	++	4
<i>Lindera pinnespora</i> NRRL 2237	-	7
<i>Penicillium chrysogenum</i> ATCC 10002	+	7
<i>Penicillium chrysogenum</i> ATCC 10002k	+	7
<i>Penicillium chrysogenum</i> ATCC 9480	+	5
<i>Penicillium purpureus</i> UI 193	-	7
<i>Penicillium vermiculatum</i> NRRL 1009	-	7
<i>Rhizopus nigricans</i> NRRL1477	-	7
<i>Rhizopus</i> species 36060	-	7
<i>Rhodotorula rubra</i> NRRL n1592	+	6
<i>Saccharomyces cerevisiae</i> (Baker's yeast)	+	5
<i>Fusarium solani</i>	-	7

-No metabolites formed. + Trace conversion of coumarin into metabolites. ++Intermediate conversion of coumarin into metabolites. +++ Maximum conversion of coumarin into metabolites.

recently reported by Nigam et al. (2013). The present study is focused on the application of microorganisms as models to study and predict fate of the investigated coumarins in mammalian systems.

MATERIALS AND METHODS

Substrates and microorganisms

Substrates were obtained from the following sources: coumarin (I) was obtained from Cairo Company (Egypt). Warfarin (III) and dicoumarol (IV) were purchased from Sigma Chemical Company Inc. St. Lewis, Mo., while umbelliferone (II) was purchased from Fluka Chemical Company Inc.; Milwaukee, Uli. Sabouraud dextrose agar was obtained from Oxoid Ltd., England. Microorganisms were obtained from American Type Culture collection (ATCC, Rockville, Maryland) or from Northern Regional Research Laboratories (NRRL, Peoria, Illinois) (Siegers and Bostelmann, 1993). A list of microorganisms is shown in Table 1.

Microbial biotransformation protocol

Preparation of the culture medium

The fermentation experiments were performed using a medium

consisting of 5 g peptone, 10 g dextrose, 10 ml glycerol, 5 g yeast extract, 5 g NaCl, 5 g K₂HPO₄ and volume adjusted to 1000 ml with distilled water. The pH was adjusted to 6.0 before sterilization at 121°C for 15 min (Myers et al., 1994).

Cultivation of microorganisms

The cultures were initiated to grow from lyophilized state by adding 1 ml of sterile water to microorganism vials. Few drops of suspended cells were streaked on Sabouraud-Dextrose plates to check the purity. The pure cultures were maintained on slants of either Potato-Dextrose agar or Sabouraud-Dextrose agar. The fresh slants were incubated for few days at room temperature before storage in a refrigerator at (4°C). The fermentation liquid cultures were initiated by transferring a few drops of suspended cells or a loopful into 25 ml sterile liquid medium present in 125 ml Erlenmeyer flask then placing on a gyratory shaker operating at 250 rpm and 27°C. The temperature was maintained at 27°C (Myers et al., 1994).

Biotransformation by *C. elegans* NRRL 1392

Fermentation of coumarin (I) was investigated using a battery of twenty-five microorganisms listed in Table 1. Stage one culture of

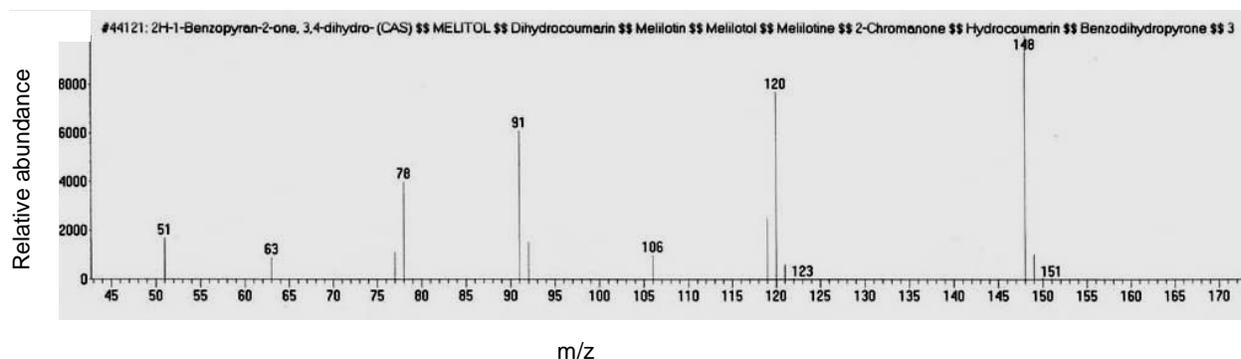


Figure 1. GC/MS spectrum of metabolite V and authentic 3, 4-dihydrocoumarin.

each investigated microorganism was initiated separately from two-week old slants (under aseptic conditions) by transferring microbial cells into 250 ml Erlenmeyer flasks, containing 50 ml of sterile liquid medium and allowed to grow for 72 h at 27°C on a gyratory shaker. Stage two cultures were obtained by transferring 5 ml of stage one culture to each of 250 ml flasks containing 50 ml of fresh liquid medium. Both substrate and organism controls were treated in the same way. The culture was allowed to grow for 24 h, before the addition of the substrate solution (0.2 mg coumarin (I) ml⁻¹ dissolved in acetone). Samples (5 ml) were periodically withdrawn from each culture and extracted successively with chloroform, then with ethyl acetate. Each extract was separately evaporated to dryness under reduced pressure and kept for further analysis. The obtained residues were dissolved in few drops of a suitable organic solvent and chromatographed on silica gel G plates (E. Merck, F 254) using solvent system i: chloroform-methanol (9.5:0.5, v/v), solvent system ii: chloroform-methanol (9:1), solvent system iii: chloroform-ethyl acetate-formic acid (4:1:0.5, v/v), solvent system iv: toluene-ethyl acetate (2:1, v/v). The plates were dried and visualized under UV light at 254 and 366 nm before and after spraying with alcoholic KOH (Feuer et al., 1976) and anisaldehyde/sulfuric reagents (Thornes et al., 1989). The results of the initial screening and the extent of biotransformation are shown in Table 1.

The same procedure was used in large scale fermentation production of each investigated coumarins into different metabolites using *C. elegans* NRRL 1392 only (Siegers and Bostelmann, 1993).

Isolation, purification and identification of metabolites

The metabolites were isolated from fermentation residues by different chromatographic methods. The residues were crystallized, separately, using the appropriate solvent. The identification of the isolated pure compounds was achieved using "Galenkamp type" melting point apparatus and different spectroscopic methods. Thermo-Finnegan mass spectrometer and API 3200TM LC/MS/MS were used. Both ¹H and ¹³C NMR spectra were taken on JEOL JNM-ECA NMR spectrometer operating at 500 MHz (¹H NMR) and 125 MHz (¹³C NMR), respectively. The chemical shift values were reported as ppm using tetramethylsilane (TMS) as an internal standard and coupling constants were expressed in Hz. Negative mode electrospray ionization mass spectra.

Large scale fermentation of coumarin (I) by *C. elegans* NRRL 1392 and isolation of metabolites

Coumarin (I, 1 g of standard coumarin) dissolved in acetone (25 ml), was equally distributed amongst 50 to 500 ml-flasks containing

stage two cultures of *C. elegans* as mentioned previously. After 4 days, the fermentation was stopped and the cells were removed by filtration.

The fermentation broth was extracted twice with an equal volume of chloroform and the solvent was distilled off under reduced pressure to give 500 mg residue. The obtained residue was chromatographed on silica gel column (20 g, 25 cm L × 1 cm D) using gradient elution method starting with hexane, hexane-chloroform, chloroform then chloroform-methanol with increasing proportions of methanol till 15%. Seventy two fractions (10 ml each) were collected. The fractions (13 to 16) eluted with hexane-chloroform (7:3) were combined, evaporated till dryness to give a pure metabolite (V) (6 mg, in yield of 0.6%). Metabolite (V) was soluble in chloroform and ether. It was subjected to GC/MS analysis as the result was represented by Figure 1. However, fractions (25 to 33) eluted with chloroform-methanol (99:1, v/v) provided 213 mg of pure metabolite (II) in 21% yield. Metabolite (II) was co-chromatographed with an authentic umbelliferone for identification.

Fractions (41 to 50) eluted with chloroform-methanol (95:5, v/v) were pooled and evaporated to a residue, which was recrystallized from methanol to give a white monoclinic crystals of metabolite (VI) (87 mg), in 9% yield. The ¹H and ¹³C NMR data of metabolite (VI) was shown in Table 4.

Large scale fermentation of umbelliferone (II) by *C. elegans* NRRL 1392

One gram of standard umbelliferone was dissolved in acetone (25 ml) and equally distributed amongst 50 to 500 ml flasks containing stage two cultures of *C. elegans* as mentioned previously. After 5 days, the fermentation was stopped and the cells were removed by filtration. Fermentation broth was extracted twice with an equal volume of chloroform and the solvent was distilled off under reduced pressure to give 510 mg residue. The residue was chromatographed on silica gel column (22 g, 25 cm L and 1 cm D) and eluted with chloroform, chloroform-ethyl acetate in gradient elution manner till pure ethyl acetate. Fractions eluted with CHCl₃-EtOAc (7.5:2.5, v/v) were evaporated under reduced pressure till dryness and recrystallized from methanol to give 125 mg of metabolite (VII) in 12.5% yield. This metabolite was subjected to ¹H and ¹³C NMR analysis as the resultant data are shown in Table 2.

Fractions eluted with CHCl₃-EtOAc (1:1, v/v) were pooled and evaporated to dryness to give 180 mg residue. This residue was further rechromatographed on Sephadex LH₂₀ column (8 g, 10 cm L × 0.5 cm D) and eluted with methanol. Thirty fractions (2 ml each) were collected. Fractions (10 to 30) gave 100 mg, a white powder of pure metabolite (VIII) in 10% yield. This metabolite possessed ¹H and ¹³C NMR analysis, where the data are shown in Table 4.

Table 2. ^1H NMR and ^{13}C NMR of metabolites VII and X.

No.	Metabolite VII	Metabolite X	No.	Metabolite VII	Metabolite X
	500 MHz- DMSO- d_6	500 MHz- CD_3OD		125 MHz- DMSO- d_6	125 MHz- CD_3OD
H-3	6.10 (d, $J=9.2$ Hz)	5.59 (s)	C-2	161.3	164.7
H-4	7.80 (d, $J=9.2$ Hz)	-	C-3	112.0	90.2
H-5	6.9 (s)	7.86 (d, $J= 7.8$ Hz)	C-4	144.9	167.1
H-6	-	7.59 (m)	C-5	112.8	123.9
H-7	-	7.28 (m)	C-6	143.3	123.2
H-8	6.70 (s)	7.29 (d, $J= 7.8$ Hz)	C-7	149.0	132.7
-	-	-	C-8	103.2	116.2
-	-	-	C-9	150.8	153.4
-	-	-	C-10	111.2	115.9

Table 3. ^1H NMR and ^{13}C NMR of metabolite IX.

No.	Metabolite IX	No.	Metabolite IX
	500 MHz-- CD_3OD		125 MHz- CD_3OD
H-3	1.9 (dd, $J=25.2, 2.3$ Hz), 2.3m	C-2	99.6
H-4	3.98 m	C-3	42.4
H-7	7.58-7.56 m	C-4	35.4
H-8	7.36-7.27 m	C-4a	101.2
H-9	7.36-7.27 m	C-5	162.3
H-10	7.89 (ddd, $J=8.4, 6.1, 1.5$ Hz)	C-6a	152.7
H-11	1.6 s	C-7	116.1
H-2', 6'	6.90 (dd, $J=8.4, 2.2$ Hz)	C-8	131.7
H-3', 5'	6.60 (dd, $J=8.4, 2.2$ Hz)	C-9	123.8
H-4'	-	C-10	122.7
-	-	C-10a	115.9
-	-	C-10b	159.7
-	-	C-11	26.4
-	-	C-1`	134.4
-	-	C-2`, 6`	128.0
-	-	C-3`, 5`	114.9
-	-	C-4`	155.4

Large scale fermentation of warfarin(III) by *C. elegans* NRRL 1392

One gram of standard warfarin was dissolved in acetone (25 ml) and equally distributed amongst 50 to 500 ml flasks containing stage two cultures of *C. elegans* as mentioned previously. After 7 days, the fermentation was stopped and the cells were removed by filtration. The fermentation broth was extracted twice with an equal volume of dichloromethane and the solvent was removed under reduced pressure to give 400 mg residue. The metabolites were separated from this residue by preparative thin layer chromatography (TLC) on silica gel G254 plates using solvent system iv as a mobile phase. Zone no. 2 was separated and extracted by ethyl acetate, the solvent evaporated to produce one compound with an $R_f = 0.23$ in solvent system iv. Metabolite (IX) was produced and purified by recrystallization from ethyl acetate giving 19 mg pure compound. Its ^1H and ^{13}C NMR spectral analysis was illustrated in Table 3.

Large scale fermentation of dicoumarol (IV) by *C. elegans* NRRL 1392

One gram of standard dicoumarol was dissolved in acetone (25 ml) and equally distributed amongst 50 to 500 ml-flasks containing stage II cultures of *C. elegans* as mentioned previously. After 8 days, the fermentation was stopped and the cells were removed by filtration. The fermentation broth was extracted twice with an equal volume of n-butanol. The extract was evaporated under reduced pressure to give 500 mg residue, which was chromatographed twice on Sephadex LH₂₀ column (25 g, 25 cm \times 1 cm), eluted with methanol.

One hundred and twenty fractions (5 ml each) were collected. Fractions (86 to 120) showed one spot. The solvent was removed under reduced pressure and the residue was recrystallized from methanol giving 32 mg of metabolite (X), in 3.2% yield. The obtained pure metabolite undergoes ^1H and ^{13}C NMR analysis as the data shown in Table 2.

Table 4. ^1H NMR and ^{13}C NMR of metabolites VI and VIII.

No.	Metabolite VI 500 MHz-- CD_3OD	Metabolite VII 500 MHz-- CD_3OD	No.	Metabolite VI 125 MHz-- CD_3OD	Metabolite VII 125 MHz-- CD_3OD
H-2	7.30	7.44 (dd, $J=8.6, 2.2$ Hz)	C-1	130.1	125.9
H-3	7.36 (m)	6.63 (dd, $J=8.6, 2.1$ Hz)	C-2	127.9	129.9
H-4	7.56 (m)	-	C-3	128.7	115.5
H-5	7.36 (m)	6.63 (dd, $J=8.6, 2.1$ Hz)	C-4	127.5	159.5
H-6	7.30	7.44 (dd, $J=8.6, 2.2$ Hz)	C-5	128.7	115.5
H-7	7.63 (d, $J=16.1$ Hz)	7.6 (d, $J=16$ Hz)	C-6	127.9	129.7
H-8	6.47 (d, $J=16.1$ Hz)	6.46 (d, $J=16$ Hz)	C-7	145.0	145.6
-	-	-	C-8	118.0	141.2
-	-	-	C-9	169.0	170.0

Cytotoxic investigation of the examined coumarins and the resultant metabolites

Potential cytotoxicity of coumarins and the isolated pure metabolites was tested using SRB assay against HEPG2 tumor cell line (Shehan et al., 1990) as follows:

(1) Cells were plated in 96-multiwell plate (10^4 cells/well) for 24 h before treatment with the compounds to allow attachment of cells to the bottom of the plate.

(2) Different concentrations of each compound under test (1, 2.5, 5 and 10 $\mu\text{g/ml}$) added to the medium containing the monolayer cell in triplicate wells were prepared for each individual dose.

(3) The wells were incubated with the compound(s) for 48 h at 37°C in an atmosphere of 5% CO_2 (using CO_2 - incubator).

(4) The medium was discarded and the cells were fixed with formalin after 48 h, washed and stained with sulforhodamine B stain.

(5) Excess stain was washed with acetic acid and attached stain was recovered with tris EDTA buffer.

(6) Color intensity was measured in an ELISA reader.

(7) The relation between surviving fraction of the cells and drug concentration is plotted to get the survival curve of each tumor cell line after treatment with the specified compound.

RESULTS AND DISCUSSION

Microbial transformation screening experiments were performed on coumarin (I) using twenty five different species of microorganisms and culminated in the detection of several metabolites. The results (Table 1) illustrated that *C. elegans* NRRL 1392 was the best microorganism capable of performing maximum conversion of coumarin at the least incubation period into metabolites followed by *Gymnascella citrina* NRRL 6050 and to lesser or negative extent for the other investigated microorganisms. The previous studies showed that fungus *C. elegans* can mimic biotransformation reactions of aromatic compounds performed by mammals (Dhalwal et al., 2008). Therefore, the filamentous fungus *C. elegans* was selected as the main microbe in the remaining biotransformation experiments.

C. elegans metabolized coumarin (I) into the corresponding metabolites (II), (VI) and (V). The metabolite

(II) was obtained as white powders soluble in ethyl acetate and methanol, which had a blue fluorescence in UV light and violet fluorescence when sprayed with alcoholic KOH reagent. It exhibited $R_f=0.37$ in solvent system I, which was identical to an authentic sample of umbelliferone (Steck and Bailey, 1969). This finding was confirmed by the ESIMS analysis, which showed $[\text{M}-1]^+$ peak at m/z 161 beside a melting point of 223 to 224°C and $m.p. = 223$ to 224°C . Metabolite (II) was identified as umbelliferone.

Pure metabolite (V) was revealed as colorless crystals soluble in chloroform and ether, with a $R_f = 0.8$ in solvent system i; $m.p. = 24$ to 25°C . It displayed a dark brown fluorescence under ultra violet light and showed no color when sprayed with anisaldehyde/sulfuric acid reagent. Metabolite (v) was subjected to GC/MS analysis as the resultant data is similar to the Liberian data of the authentic sample of 3, 4-dihydrocoumarin represented by Figure 1. Metabolite (V) was identified as 3, 4-dihydrocoumarin.

White monoclinic crystals of metabolite (VI), which is soluble in methanol displayed R_f values of 0.25 and 0.61 in solvent systems i and ii, respectively. It exhibited a dark brown color when sprayed with alcoholic KOH, no color when sprayed with anisaldehyde/sulfuric acid reagent. Metabolite (VI) has a $m.p. = 133^\circ\text{C}$ and was subjected to ESIMS analysis obtaining a quasi molecular ion at $(m/z): 149 [\text{M}+1]^+$. The ^1H and ^{13}C NMR data of metabolite (VI) represented in Table 4 were in concordance with the data reported for reference transcinnamic acid (Kalinowska et al., 2007; EL-Askary et al., 2008), so metabolite (VI) was identified as trans cinnamic acid.

Umbelliferone (II) was metabolized by *C. elegans* into metabolites (VII) and (VIII). Metabolite (VII) was obtained as yellowish green powder, which exhibited R_f values of 0.17 and 0.53 in solvent systems ii and iii, respectively. It showed a dark brown fluorescence in UV light and no color when sprayed with anisaldehyde/sulfuric acid reagent. ^1H and ^{13}C NMR data of metabolite (VII) are shown in Table 2, which lacked the H-6, H-7 signals

resonating at δ_H 7.19, 7.45 in coumarin (Perel'son et al., 1970). Furthermore, H-5 signal for this metabolite underwent an upfield shift of 0.51 ppm, which is consistent with the ortho effect of introduced hydroxyl group (Sothers, 1972) at C-6. Examination of ^{13}C NMR spectral data of the metabolite (VII) (Table 2) in comparison with those of coumarin (Kumari et al., 2004) confirmed the hydroxylation at positions attached to C-6 and C-7 as the ipso carbons exhibited a downfield shift of 20 ppm, 17.3 ppm. However, C-5 (ortho carbon) experienced an upfield shift of 15 ppm and C-8 (orthocarbon) showed a upfield shift of 13.5 ppm. All the aforementioned data were identical to data reported for a esculetin (VII) (Steck and Bailey, 1969). Thus, metabolites (VII) was identified as 6,7-dihydroxycoumarin (aesculetin).

However, metabolite (VIII) is a green powder soluble in methanol and ethyl acetate, possessed R_f values of 0.28 and 0.46 in solvent systems i and ii, respectively. It has a violet fluorescence in UV light and pink color when sprayed with anisaldehyde/sulfuric acid reagent and a melting point at 209 to 211°C. The ^1H NMR and ^{13}C NMR of this metabolites (Table 4) was identical to the corresponding data of p-coumaric acid (EL-Askary et al., 2008), which was structurally elucidated as p-coumaric acid.

Warfarin (III) was biotransformed by the same microorganism into metabolite (IX), a small needle crystal soluble in methanol and ethyl acetate. It exhibited a faint brown fluorescence under UV, a faint green color with KOH and with a m.p. = 200°C. ^1H and ^{13}C NMR data of metabolite IX are shown in Tables 3, exhibiting two aromatic doublet of doublet signals resonating at δ_H 6.60, 6.90 ppm ($J=8.4, 2.2$ Hz), assigned for protons H-3',5' and H-2',6', respectively, which indicated a 1,4-disubstituted benzene ring (Rizzo and Davis, 1989), compared with warfarin (Gebauer, 2007). On the other hand, ^{13}C NMR of this metabolite (Table 3) compared to those of warfarin (Pisklak et al., 2003) showed C-4' signal strongly deshielded, while the C-3', C-5' signal is shifted upfield by 10 to 12.1 ppm. Furthermore, C-2', C-6' signal underwent a slight upfield shift of about 0.20 ppm. These results are in close agreement with substituent effect of a hydroxyl group on aromatic ring carbons (Sothers, 1972) and support the identification of the metabolite IX as 4'-hydroxy warfarin. *C. elegans* hydroxylation of warfarin (III) at 4' position was demonstrated to proceed via an arene oxide intermediate and displayed NIH shift (Pisklak et al., 2003).

Dicoumarol (IV) undergoes a specific metabolic reaction by *C. elegans* producing metabolite X as yellowish green powder, which possessed R_f value of 0.35 using solvent system ii with a dark brown fluorescence under UV light and faint green color when sprayed with anisaldehyde/sulfuric acid reagent. The metabolite was soluble in methanol and hot water. The interpretation of ^1H data of metabolite X (Table 2) showed

the absence of H-4 signals resonating at δ_H 7.76 ppm, which is consistent with the ortho effect of introduced hydroxyl group (Sothers, 1972) at C-6.7.45 and 7.76 ppm in coumarin (Perel'son et al., 1970). H-3 for metabolite X revealed an upfield shift of 0.85 ppm. ^{13}C NMR spectral data of this metabolite illustrated that C-4 exhibited a downfield shift of 26.3 ppm and C-3 (ortho carbon) showed an upfield shift of 26.3 ppm in comparison with that of coumarin (Kumari et al., 2004) confirmed the hydroxylation of C-4 position. All the aforementioned data were identical to data reported for 4-hydroxycoumarin (X) (Traven et al., 1997), thus metabolites X was identified as 4-hydroxycoumarin.

The course of studying the metabolism of the investigated coumarins by *C. elegans* illustrated that several reaction was attempted. It hydroxylated coumarin (I), umbelliferone (II) and warfarin (III) into the corresponding hydroxyl derivatives, such as umbelliferone (II), 6, 7-dihydroxycoumarin (VII) and 4'-hydroxywarfarin (IX), respectively (Figure 2). Aromatic hydroxylation of coumarin (I) and umbelliferone (II) at positions 7,6 by *C. elegans* has also a parallel in the mammalian systems (Lake, 1999; Shimada et al., 1996).

Other reduction reaction of coumarin (I) by *C. elegans* was attempted producing 3, 4-dihydrocoumarin (V), which was confirmed by GC-MS analysis. Formation of 3, 4-dihydroderivative takes place in coumarin (I) only, which indicated that the enzyme may have narrow substrate specificity.

Two main polar transcinamic acid (VI) and p-coumaric acid (VIII) were obtained by a cleavage reaction (opening of the lactone ring of the heterocyclic ring system) of coumarin (I) and umbelliferone (II), respectively. This reaction is limited for coumarin (I) and umbelliferone (II), suggesting its (narrow) substrate specificity as the exact mechanism of this reaction needs further investigation.

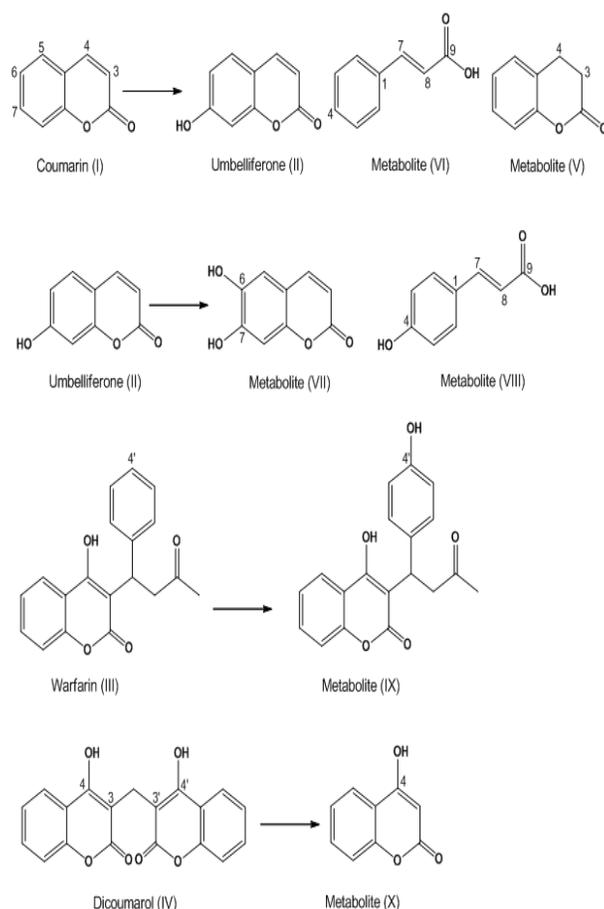
However, *C. elegans* converted dicoumarol (IV) by a specific metabolic reaction as cleavage of methylene bridge into 4-hydroxycoumarin(X), in a mechanism which awaits further investigation.

Coumarins and related derivatives are widely used in the treatment of many diseases. Thus before ending this investigation, an assay was conducted on coumarin substrates under investigation, as well as the isolated pure metabolites for their cytotoxic activity (Table 5). The cytotoxicity of all coumarins was tested against HEPG2 tumor cell line adopting SRB assay (Maucher and Von Angerar, 1994) and IC_{50} determination and compared to that of a standard doxorubicin. All tested coumarins exhibited much higher IC_{50} (low cytotoxic effect) compared to doxorubicin except for 3, 4-dihydrocoumarin, warfarin and 6, 7-dihydroxycoumarin and further studies against other cell lines are worthwhile. All biological studies revealed that the investigated coumarins were metabolized into products, which were far less active, except for 6, 7-dihydroxycoumarin (VII) and 3, 4-dihydrocoumarin (V).

Table 5. Effect of coumarins and pure metabolites on the cell viability of HepG2 cell line.

Compound	Concentration ($\mu\text{g/ml}$)				IC 50 (μg)
	1	2.5	5	10	
Coumarin	0.71	0.48	0.29.	0.29	2.55
Umbelliferone	0.75	0.58	0.32	0.23	3.02
Dicoumarol	0.81	0.42	0.31	0.22	2.28
Warfarin	0.39	0.36	0.36	0.22	0.67
trans-Cinnamic acid	0.85	0.68	0.39	0.26	3.83
3,4-Dihydrocoumarin	0.23	0.24	0.33	0.43	0.65
6,7-Dihydroxycoumarin	0.35	0.34	0.33	0.32	0.75
p-coumaric acid	0.80	0.61	0.52	0.41	5.03
4-Hydroxycoumarin	0.75	0.52	0.41	0.27	2.89
4'-Hydroxywarfarin	0.92	0.73	0.53	0.45	5.44
Doxorubicin*	0.34	-	0.35	0.34	0.60

*A standard antitumor compound

**Figure 2.** Metabolism of coumarins by *C. elegans*.

Conclusion

The fungus *C. elegans* NRRL 1392 metabolized coumarin (I) into umbelliferone (II), 3, 4-dihydrocoumarin

(V) and trans cinnamic acid (VI). Umbelliferone was also biotransformed into 6, 7-dihydroxycoumarin (VII) and p-coumaric acid (VIII). On the other hand, warfarin (III) was biotransformed into 4'-hydroxywarfarin (IX) and also

dicoumarol (IV) was transformed into 4-hydroxycoumarin (X) by the same microorganism. The cytotoxic effect of coumarin substrates and their metabolites was investigated revealing that both 6, 7-dihydroxycoumarin (VII) and 3, 4-dihydrocoumarin (V) were more active than a standard doxorubicin (positive control).

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