

Microbial Transformations of 3-methoxyflavone by Strains of *Aspergillus niger*

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Abstract

Microbial transformation of 3-methoxyflavone into 3'-hydroxyflavon-3-yloxymethyl myristate was presented. Six filamentous fungi were used as biocatalysts: a wild strain of *Aspergillus niger* KB, its four UV mutants (*A. niger* MB, SBP, SBJ, 13/5) and the strain of *Penicillium chermesinum* 113. The highest yields were observed for the strains of *A. niger* KB and *A. niger* SBP (69.8% and 63.1%, respectively).

Key words: *Aspergillus niger*, *Penicillium chermesinum*, biotransformation, 3-methoxyflavone, myristic acid

Microbial transformation of flavonoid compounds is a natural method which may be a green alternative to chemical synthesis. Using biotransformation we can modify structures of compounds in order to improve their biological properties and to increase their hydrophilicity or bioaccessibility (Das and Rosazza, 2006; Wang *et al.*, 2010).

Regioselective *O*-demethylation of tangeretin and 3-hydroxytangeretin by *Aspergillus niger* gave 4'-*O*-demethylated biotransformation products (Buisson *et al.*, 2007) while transformations of 7,8-dimethoxyflavone by *Mucor ramannianus* gave five biotransformation products involving hydroxylation at C-3' and C-4' with the methoxyl groups retained and products of hydroxylation in the B-ring along with demethylation at C-7 or/and C-8 (Herath *et al.*, 2009).

This manuscript reports the microbial demethylation of 3-methoxyflavone by strains of *Aspergillus* and *Penicillium*, followed by non-typical esterification with a fatty acid and hydroxylation in the B-ring.

The analytical procedures were as described previously (Kostrzeva-Susłow and Janeczko, 2012a; 2012b).

The substrate for biotransformation – 3-methoxyflavone was purchased from Sigma-Aldrich, Poznań, Poland.

3-Methoxyflavone (C₃₀H₃₈O₆): M.p. 114–115°C. Rt 19.35 min (HPLC).

¹H NMR (DMSO-d₆) δ: 3.81 (3H, s, -OCH₃), 7.49 (1H, t, J_{6,5} = 8.0, J_{6,7} = 7.0 Hz, H-6), 7.58 (3H, m, H-3', H-4', H-5'), 7.75 (1H, d, J_{8,7} = 8.2 Hz, H-8), 7.82 (1H, ddd, J_{7,5} = 1.5, J_{7,6} = 7.0, J_{7,8} = 8.2 Hz, H-7), 8.05 (2H, m, H-2', H-6'),

8.10 (1H, dd, J_{5,6} = 8.0, and J_{5,7} = 1.5 Hz, H-5); ¹³C NMR (DMSO-d₆) δ: 60.2 (-OCH₃), 119.0 (C-8), 124.1 (C-10), 125.4 (C-6), 125.6 (C-5), 128.8 (C-2', C-6'), 129.2 (C-3', C-5'), 131.0 (C-4'), 131.4 (C-1'), 134.6 (C-7), 141.3 (C-3), 155.3 (C-2), 155.5 (C-9), and 174.4 (C-4).

In this research we used a wild strain of *A. niger* KB and four UV mutants of *A. niger* (13/5, SBJ, SBP, MB). The KB strain comes from the collection of the Department of Biotechnology and Food Microbiology of Wrocław University of Environmental and Life Sciences (Poland), strain 13/5 comes from the University of Life Sciences in Lublin (Poland) and strains SBJ, SBP and MB come from Wrocław University of Economics (Poland). Microorganisms were maintained on sterilized potato slants at 5°C. The wild strain of *P. chermesinum* 113 was obtained from the collection of the Department of Chemistry of Wrocław University of Environmental and Life Sciences (Poland). The microorganism was maintained on Sabouraud 4% dextrose-agar slopes and freshly subcultured before use in the transformation experiments.

Screening tests and preparative-scale biotransformations were carried out according to the procedure described earlier (Kostrzeva-Susłow and Janeczko, 2012a; 2012b). The spectral data of the product obtained are presented below.

3'-Hydroxyflavon-3-yloxymethyl myristate (C₃₀H₃₈O₆): Yellow oily liquid. Rt 17.06 min (HPLC). Purity 97% (HPLC). HRESI-MS: *m/z* = 495.1032 [M+H⁺]; found 495.1026. ¹H NMR (DMSO-d₆) δ: 0.86 (3H, t, J = 7.0 Hz, 13''-CH₃), 1.229–1.231 (14H, m, H-4''-H-10''),

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1.239–1.261 (6H, m, H-3",11",12"), 1.55 (2H, tt, $J=7.67$ Hz, H-2"), 2.25 (2H, t, $J=7.4$ Hz, H-1"), 5.22 (2H, s, O-CH₂-O), 7.30 (1H, m, H-4'), 7.39 (1H, ddd, $J_{6,5}=7.9$, $J_{6,7}=7.1$ Hz, $J_{6,8}=1.1$ Hz, H-6), 7.49 (3H, m, H-8, 2', 5'), 7.66 (1H, ddd, $J_{7,8}=8.4$ Hz, $J_{7,6}=7.1$ Hz, $J_{7,5}=1.7$ Hz, H-7), 7.85 (1H, m, H-6'), 7.91 (1H, s, -OH), and 8.19 (1H, dd, $J_{5,6}=8.0$, $J_{5,7}=1.7$ Hz, H-5). ¹³C NMR (DMSO-d₆) δ: 14.3 (C-13"), 25.7 (C-2"), 29.1 (C-10"), 29.3 (C-8"), 29.3 (C-9"), 29.4 (C-7"), 29.5 (C-6"), 29.6 (C-5"), 29.7 (C-4"), 29.8 (C-3"), 30.1 (C-11"), 31.8 (C-12"), 34.1 (C-1'), 83.0 (O-CH₂-O), 111.5 (C-2'), 116.9 (C-4'), 117.3 (C-6'), 119.4 (C-8), 124.2 (C-10), 125.6 (C-6), 125.7 (C-5), 130.0 (C-5'), 132.6 (C-1'), 134.7 (C-7), 149.2 (C-3), 151.0 (C-3'), 155.5 (C-2), 155.7 (C-9), 175.1 (C-4), and 180.8 (O=C=O).

Screening tests revealed six strains of filamentous fungi capable of biotransformation of 3-methoxyfla-

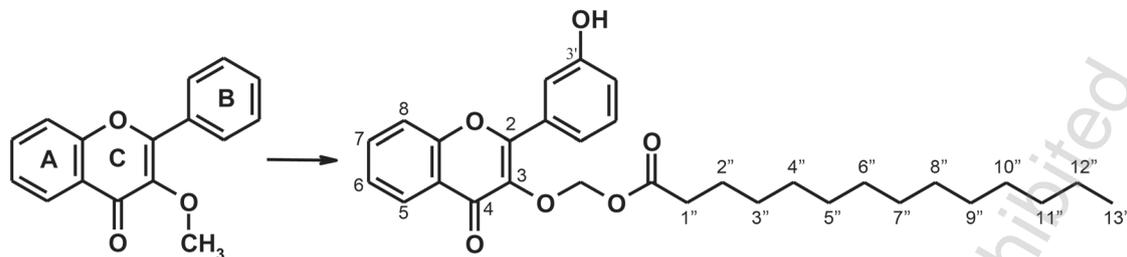
vone: a wild strain of *A. niger* KB, its four UV-mutants (*A. niger* MB, SBP, SBJ, 13/5) and the strain of *P. chermesinum* 113. The initial study was performed according to two procedures: in the first the substrate was added to the cultivation media at the time of inoculation with the microorganism, whereas in the second, it was added 72 hours after the inoculation. After 3, 6 and 9 days samples were analyzed for product and unreacted substrate (Table I, II). All of the tested strains of *Aspergillus* transformed 3-methoxyflavone into 3'-hydroxyflavon-3-yloxymethyl myristate, using either cultivation procedure. *P. chermesinum* 113 was also capable of this transformation, but only when the substrate was added to the cultivation media in the last phase of logarithmic growth of the mycelium (Table II). For all of the tested *Aspergillus* strains the biotransformation was more efficient when the substrate was added to the cultivation

Table I
Biotransformation of 3-methoxyflavone (substrate added at the time of inoculation)
– yield (%) of product determined by HPLC

Micro-organism	Time of incubation (days) →	3	6	9
	Product	Yield (%)		
<i>A. niger</i> KB	3'-hydroxyflavon-3-yloxymethyl myristate	53.7	64.8	69.8
	Unreacted substrate	38.2	24.1	15.5
<i>A. niger</i> MB	3'-hydroxyflavon-3-yloxymethyl myristate	25.1	30.7	37.2
	Unreacted substrate	73.2	63.2	53.0
<i>A. niger</i> SBP	3'-hydroxyflavon-3-yloxymethyl myristate	38.7	50.4	63.1
	Unreacted substrate	60.0	46.3	27.3
<i>A. niger</i> SBJ	3'-hydroxyflavon-3-yloxymethyl myristate	22.5	23.7	24.9
	Unreacted substrate	75.9	70.9	59.8
<i>A. niger</i> 13/5	3'-hydroxyflavon-3-yloxymethyl myristate	10.1	20.3	20.5
	Unreacted substrate	87.8	73.2	70.0

Table II
Biotransformation of 3-methoxyflavone (substrate added 72 hours after inoculation)
– yield (%) of product determined by HPLC

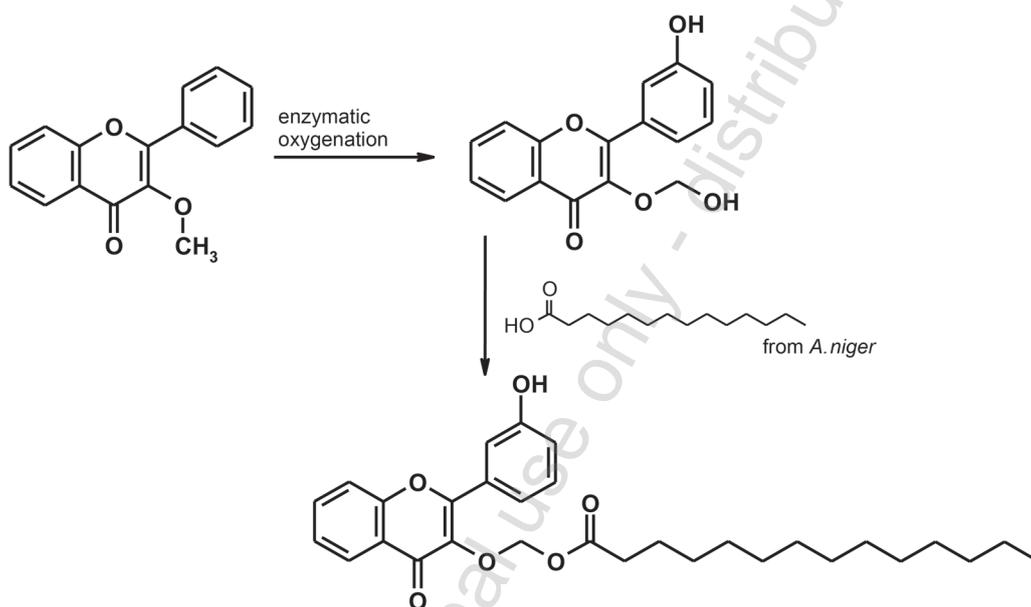
Microorganism	Time of incubation (days) →	3	6	9
	Product	Yield (%)		
<i>A. niger</i> KB	3'-hydroxyflavon-3-yloxymethyl myristate	13.7	16.2	18.4
	Unreacted substrate	81.2	70.7	65.0
<i>A. niger</i> MB	3'-hydroxyflavon-3-yloxymethyl myristate	16.3	16.2	16.2
	Unreacted substrate	80.1	80.0	78.2
<i>A. niger</i> SBP	3'-hydroxyflavon-3-yloxymethyl myristate	26.2	26.9	27.3
	Unreacted substrate	72.1	69.7	65.1
<i>A. niger</i> SBJ	3'-hydroxyflavon-3-yloxymethyl myristate	13.3	15.6	17.2
	Unreacted substrate	80.4	75.1	70.1
<i>A. niger</i> 13/5	3'-hydroxyflavon-3-yloxymethyl myristate	12.7	13.5	14.1
	Unreacted substrate	80.3	77.0	70.2
<i>P. chermesinum</i> 113	3'-hydroxyflavon-3-yloxymethyl myristate	16.7	19.2	23.8
	Unreacted substrate	80.1	73.3	65.1



Scheme 1. Biotransformation of 3-methoxyflavone in the culture of *A. niger* (KB, MB, SBP, SBJ, 13/5) and *P. chermesinum* 113.



Scheme 2. Mechanism of demethylation.



Scheme 3. Probable course of biotransformation of 3-methoxyflavone.

media at the time of inoculation with the microorganism. *A. niger* KB and *A. niger* SBP gave the highest yields of the product after 9 days, *i.e.* 69.8% and 63.1%, respectively. The least efficient strain for this biotransformation was *A. niger* 13/5 (Table I, II).

The preparative scale biotransformation of 3-methoxyflavone was carried out using the strain of *A. niger* KB in a 9-day reaction. 3'-Hydroxyflavon-3-yl myristate was isolated in 65% yield (Scheme 1) and identified by ^1H NMR and ^{13}C NMR. In the ^1H NMR spectrum the signal of the substrate methoxyl group at C-3 ($\delta = 3.81$ ppm) disappeared and a two proton singlet at $\delta = 5.22$ ppm attributed to a $-\text{O}-\text{CH}_2-\text{O}-$ group appeared. Seven new signals integrating for a total of 24 protons and a singlet of 3H at $\delta = 0.86$ ppm indicate the presence of a 13-carbon alkyl chain of a saturated fatty acid (myristic acid) bonded at C-3 of the flavone by means of the ester bond. The ester carbonyl group is confirmed by the signal at $\delta = 180.8$ ppm in

the ^{13}C NMR. The presence of a hydroxyl group in the B-ring was proved by the new one-proton singlet at $\delta = 7.91$ ppm which was not present in the spectrum of the substrate. The location of the hydroxyl group was determined by analyzing the shape and chemical shift of the B-ring protons. The 2' and 6' protons and the 4' and 5' protons are nonequivalent in the ^1H NMR of the product. In the ^{13}C NMR the C-3' signal is shifted from $\delta = 129.2$ ppm for the substrate to $\delta = 151.0$ ppm for the product.

The mechanism of oxidative *O*-demethylation of methyl ethers by cytochrome P450 proposed by Watanabe is presented in Scheme 2 (Watanabe *et al.*, 1982).

In the biotransformation of 3-methoxyflavone catalyzed by the strains of *Aspergillus* and *Penicillium* the intermediate hemiacetal is presumably esterified with myristic acid present in the cultivation mixture. Myristic acid is produced by the strains of *Aspergillus*. The presence of considerable myristic acid has been observed

in conidiophores of *A. flavus* (Budínská *et al.*, 1981) in the cells of *A. ochraceus* (Chavant and Sandolle, 1977) and in the cells of *A. niger* (Parang *et al.*, 1996).

The first step of enzymatic oxidation of 3-methoxyflavone resembles metabolism of methoxy derivatives in mammals. The reaction is probably catalyzed by the fungal monooxygenases of cytochrome P-450. The proposed mechanism of the biotransformation is presented in Scheme 3. Earlier research on microbial transformations of monosubstituted flavones with methoxyl groups in the A-ring (Kostrzewa-Susłow *et al.*, 2012) indicate that the strains of *Aspergillus* and *P. chermesinum* used for this study perform non-typical demethylation which ends with introduction of myristic acid in the favourable C-3 flavone position.

Conclusions. Transformation of 3-methoxyflavone in the cultures of *A. niger* (KB, MB, SBP, SBJ, 13/5) and *P. chermesinum* 113 is a two-step process involving enzymatic oxidation of both the methoxyl group and the B-ring of the flavone and then esterification of the $-O-CH_2-OH$ group with myristic acid. The highest yield of transformation was achieved for the strain *A. niger* KB, when the substrate was added at the time of inoculation.

Acknowledgements

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