

HHS Public Access

Author manuscript *J Nat Prod.* Author manuscript; available in PMC 2018 February 21.

Published in final edited form as:

J Nat Prod. 2016 February 26; 79(2): 447–450. doi:10.1021/acs.jnatprod.5b00972.

N-acyl dehydrotyrosines, tyrosinase inhibitors from the marine bacterium *Thalassotalea* sp. PP2-459

Robert W. Deering^{†,⊥}, Jianwei Chen^{†,‡,⊥}, Jiadong Sun[†], Hang Ma[†], Javier Dubert[§], Juan L. Barja[§], Navindra P. Seeram[†], Hong Wang[‡], and David C. Rowley^{†,*}

[†]Department of Biomedical and Pharmaceutical Sciences, College of Pharmacy, University of Rhode Island, Kingston, RI, United States

[‡]College of Pharmaceutical Science, Zhejiang University of Technology, Hangzhou 310014, China

[§]Department of Microbiology and Parasitology, CIBUS- Faculty of Biology and Aquaculture Institute, University of Santiago de Compostela, Santiago de Compostela 17582, Spain

Abstract

Thalassotalic acids A–C and thalassotalamides A and B are new N-acyl dehydrotyrosine derivatives produced by *Thalassotalea* sp. PP2–459, a Gram-negative bacterium isolated from a marine bivalve aquaculture facility. The structures were elucidated via a combination of spectroscopic analyses emphasizing two-dimensional NMR and high-resolution mass spectral data. Thalassotalic acid A (1) displays *in vitro* inhibition of the enzyme tyrosinase with an IC₅₀ value (130 μ M) that compares favorably to the commercially-used control compounds kojic acid (46 μ M) and arbutin (100 μ M). These are the first natural products reported from a bacterium belonging to the genus *Thalassotalea*.

Graphical Abstract



Marine bacteria continue to be a highly productive source of new natural products with 408 new compounds reported from 2011–2013.^{1–3} While marine Gram-positive bacteria, especially actinomycetes, have garnered the most attention to date, genome mining and

Corresponding Author: drowley@uri.edu; Tel: 401-874-9228.

¹Author Contributions

R. W. Deering and J. Chen contributed equally to this work.

Supporting Information Available: 1D- and 2D-NMR and HRESI mass spectra for compounds 1–5. The Supporting Information is available free of charge on the ACS Publications website at DOI:

isolation-driven discovery efforts have raised awareness for the biosynthetic potential of marine Gram-negative bacteria (GNB).^{4, 5} Recognition that marine GNB produce such molecules as the didemnins,^{6, 7} and quite likely the bryostatins,⁸ highlights the potential for discovery of molecules with biomedical value. The marine γ -proteobacteria have been a particularly productive taxonomic group of GNB for natural product studies, yielding notable antibiotics such as andrimid and the moiramides, as well as the signaling molecule AI-2.^{9, 10} These studies and others portend the future success of discovery efforts aimed at marine GNB.

Thalassotalea is a new Gram-negative, aerobic, γ -proteobacteria genus.¹¹ This genus was first characterized in 2014 following reclassification of members of the genus *Thalassomonas*.¹¹ There have been only 10 species of *Thalassotalea* discussed in the literature and they have been primarily studied for their agarolytic properties, commensalism with marine animals, and pathogenic contribution to coral diseases.^{11–16} To date, there have been no published secondary metabolite studies on this genus.^{11–14}

The enzyme tyrosinase is responsible for three transformations in the conversion of tyrosine to melanin, which is responsible for pigmentation of skin, hair, and eyes.¹⁷ It specifically catalyzes the conversion of L-tyrosine to 3,4-dihydroxyphenylalanine (L-DOPA), L-DOPA to dopaquinone, and 5,6-dihydroxyindole (DHI) to indole-quinone.¹⁷ Because of tyrosinase's role in pigmentary diseases such as albinism and melasma, as well as cosmetic relevance to increase or decrease skin or hair color and pigmentation, compounds that can modulate tyrosinase have potential therapeutic and cosmetic applications.¹⁸ Additionally, tyrosinase is responsible for enzymatic browning of foods and inhibition of this process could lead to improved agricultural lifespans and nutrition.¹⁹

In this paper, we report new *N*-acyl dehydrotyrosine derivatives (1–5) produced during laboratory cultivation of *Thalassotalea* sp. PP2-459, formerly classified as *Thalassomonas* sp. PP2-459, a strain isolated from a carpet-shell clam (*Ruditapes decussata*) harvested in a bivalve hatchery located in Galicia (NW Spain).²⁰ PP2-459 was previously determined to have quorum quenching ability against key bivalve aquaculture pathogens, including *Vibrio anguillarum*.²⁰ Hence, PP2-459 has been proposed as a candidate for probiotic applications in aquaculture. Due to structural similarity with known tyrosinase inhibitors such as *N*-acylated tyrosines, 1–5 were evaluated for their inhibition of the tyrosinase enzyme.

Thalassotalea PP2-459 was cultured in YP seawater medium at 25 °C shaking and 200 rpm for two days. The cultures were centrifuged at 10,000 g for 10 min to substantially remove the cells, and the resulting supernatants were extracted with ethyl acetate. Purification efforts using reversed-phase (C18) chromatography methods yielded new compounds 1-5.

Thalassotalic acid A (1) was isolated as a light brown powder. The molecular formula was determined to be $C_{19}H_{27}NO_4$ using HRESIMS (m/z 332.1866 [M-H]⁻, calcd for $C_{19}H_{26}NO_4$, 332.1867), indicating 7 degrees of unsaturation. The presence of a decanoyl chain was supported by ¹H-NMR resonances at δ_H 0.90 (t, 3H, H-10'), 1.29–1.31 (m, 12H, H-4'-H-9'), 1.59 (m, 2H, H-3'), and 2.28 (t, 2H, H-2') that comprised a spin system in the ¹H-¹H COSY spectrum. ¹³C-NMR resonances at δ_C 172.0 (C-1') and 166.7 (C-1) indicated

the presence of two carbonyl carbons, the former belonging to the decanoyl chain based on HMBC correlations from H-2' and H-3'. In the aromatic region, two doublets at $\delta_{\rm H}$ 6.80 (H-6, J= 8.7 Hz) and 7.51 (H-5, J= 8.7 Hz) provided evidence for a para-substituted benzene ring. A HSQC correlation assigned a sharp singlet at $\delta_{\rm H}$ 7.22 (H-3) as an olefinic methine attached to carbon at $\delta_{\rm C}$ 132.2. HMBC correlations from H-5 to C-3 and H-3 to C-5 indicated that the C-3 methine was the benzylic position of the para-substituted aromatic ring. Consideration of the molecular formula and further HMBC correlations from the C-2-NH to C-1, C-2, and C-3, as well as correlations from H-3 to C-1 and C-2, established the α - β unsaturated tyrosine (Fig. 1). The (Z)-configuration of the double bond was determined based on NOE correlations shown in Fig. 1. TOF-MS/MS experiments further corroborated the proposed structure. Product ions were observed indicating the loss of CO₂H (m/z 288.2, [M-45]), the decanoyl chain (m/z 178.0, [M-155]), and both the carboxylic acid and the acyl chain (m/z 134.0, [M-199]).

The structure of thalassotalic acid B (2) was determined by comparison to spectroscopic data of 1. A molecular ion at m/z 332.1872 [M-H]⁻ indicated the same molecular formula as 1: $C_{19}H_{27}NO_4$. A key difference in the ¹H-NMR spectrum was the presence of a sharp doublet at δ_H 0.90 that integrated for 6 protons, suggesting a terminal branch in the aliphatic chain. Analysis of the HSQC spectrum revealed an aliphatic methine at δ_H 1.55 and δ_C 27.4, whose ¹H-NMR resonance coupled to the C-8' methyl groups in the COSY spectrum, thus confirming the terminal (CH₃)₂CH- group. The remaining structure was determined to be identical to 1 based on comparison of ¹³C and 2D NMR data and TOF-MS/MS evidence.

Thalassotalic acid C (3) displayed a HRESIMS ion $(m/z 318.1699 [M-H]^-$, calcd for $C_{18}H_{24}NO_4$, 318.1711) consistent with a molecular formula of $C_{18}H_{25}NO_4$, which is one CH₂ unit fewer than 1 and 2. Comparison of ¹H- and ¹³C-NMR data with 1 and 2 revealed that 3 shared the same core tyrosine structure, but incorporated an unbranched chain containing one fewer methylene unit than 1. Thus, 3 was determined to be the nonanoyl analog of 1. TOF-MS/MS further corroborated this structure with the characteristic losses of CO₂H (*m/z* 274.2, [M-45]), the nonanoyl chain (*m/z* 178.1, [M-141]), and both the carboxylic acid and the acyl chain (*m/z* 134.1 [M-185]).

Thalassotalamide A (4) was isolated as an amorphous brown powder. HRESIMS indicated that the molecular formula was $C_{19}H_{28}N_2O_3$ based on a molecular ion (*m/z* 331.2021 [M-H] ⁻, calcd for $C_{19}H_{27}N_2O_3$, 331.2027). Thus, 4 had an additional NH and one less O than 1. Examination of the ¹H-NMR spectrum of 4 indicated strong similarity to 1 based on resonances attributed to the *p*-substituted benzene ring (two doublets each integrating as 2H at δ_H 6.78 and 7.42) and unbranched aliphatic chain. The outstanding difference was two broad singlets at δ_H 7.05 and 7.28, indicating that 4 was likely the amide analog of 1. TOF-MS/MS confirmed this suspicion as fragment ions of *m/z* 288.2 [M-44] and *m/z* 134.1 [M-198] were observed, indicating that the α - β unsaturated carbonyl was an amide and not a carboxylic acid. Analysis of ¹³C- and 2D-NMR and further confirmed the structure of 4.

The structure of thalassotalamide B (5) was elucidated by comparison to spectral data of 2 and 4. HRESIMS indicated that the molecular formula of 5 was the same as 4 $(C_{19}H_{28}N_2O_3)$ based on a molecular ion (*m/z* 331.2015 [M-H]⁻, calcd for $C_{19}H_{27}N_2O_3$,

331.2027). In the ¹H-NMR spectrum, a doublet at $\delta_{\rm H}$ 0.89 integrating 6H suggested that **5** had the same branched aliphatic chain as **2**. Comparison of ¹³C-NMR data from the aliphatic chain of **2** to **5** confirmed they had the same 8-methyl nonanoyl aliphatic chain. Comparison of the remaining ¹³C-NMR data to those of **4**, as well as the prominent loss of the amide in TOF-MS/MS (*m/z* 288.2), indicated that **5** was the amide derivative of **2**.

N-acyl tyrosines have previously been studied for their effects on pigmentation of skin and hair.^{21–24} These compounds have been implicated in both increasing and decreasing pigmentation.^{21–24} Because of the similarity in structure to *N*-acyl tyrosines, we evaluated compounds **1–5** for their effects on tyrosinase enzyme activity compared to the positive controls arbutin (**6**) and kojic acid (**7**) that have been widely used in skin whitening products. ^{25–27} Compounds **1**, **2**, and **3** exhibited tyrosinase enzyme inhibition with IC₅₀ values of 130 \pm 10 µM, 470 \pm 10 µM, and 280 \pm 10 µM, respectively (Table 2). The IC₅₀ value for **1** compared favorably to the **6** (100 \pm 10 µM), but was less effective than **7** (IC₅₀ = 46 \pm 2 µM). Compounds **4** and **5** were inactive at the maximum tested concentration (1 mM). These data support the requirement for a carboxylic acid and straight aliphatic chain for increasing enzyme inhibition within this structural class of tyrosinase inhibitors. While these N-acyl dehydrotyrosine compounds are not as potent as **6** or **7**, **1** displays comparable tyrosinase inhibition and could be useful as a whitening agent or as an agricultural additive to prevent the browning of foods. ¹⁹



N-acyl derivatives of amino acids, including tyrosine, have been previously produced via heterologous expression of environmental DNA in *Escherichia coli*. These molecules are products of *N*-acyl amino acid synthases, which have been observed as single genes or as part of more complex biosynthetic sequences.^{28–30} While the compounds discussed here are structurally similar, the dehydrogenation across the α - β carbons of the tyrosine, the terminal amide functional group in **4** and **5**, and the branched acyl chains in **2** and **5** represent biosynthetic variations not present in heterologously produced *N*-acyl tyrosines.²⁹ Additionally, these are the first natural products reported from a bacterium belonging to the genus *Thalassotalea*.

EXPERIMENTAL SECTION

General Experimental Procedures

All chemical reagents and solvents were purchased from Sigma-Aldrich (St. Louis, MO) and acquired via Wilkem Scientific (Pawcatuck, RI). NMR experiments were conducted using an Agilent NMRS 500 MHz (Agilent Technologies, Santa Clara, CA) with $(CD_3)_2SO$ as the solvent (referenced to residual DMSO at $\delta_H 2.54$ and $\delta_C 39.5$) at 25 °C. Electrospray ionization mass spectra (ESIMS) were acquired using an AB Sciex TripleTOF® 4600 spectrometer (Framingham, MA) in the negative ion mode. Flash chromatography was completed with a Combiflash Rf200 equipped with an 86 g C₁₈ RediSep®Rf High Performance Gold column (Teledyne ISCO, Lincoln, NE). HPLC experiments were performed on a Hitachi Elite LaChrom system (Pleasanton, CA) equipped with a diode array detector (DAD) model L-2450, pump L-2130, and autosampler L-2200. Semi-preparative HPLC experiments were completed with a Waters XBridgeTM Prep C18 5 µm, 10 × 250 mm column (Waters Corporation, Milford, MA).

Bacterial Strain and Culture Conditions

The bacteria strain (PP2-459) was isolated from a bivalve hatchery in Galicia, Spain and identified as a *Thalassotalea* sp. based on 16S rRNA sequence comparison (European Nucleotide Archive; accession number LN898116).²⁰ It has been deposited in the Spanish Type Culture Collection (CECT) under restricted access. Cultures were maintained in YP seawater medium (1 g yeast extract, 5 g peptone, 22.5 g Instant Ocean (Spectrum Brands, Blacksburg, VA) per liter of deionized water) at 25 °C and shaking at 200 rpm.

Extraction and Isolation

A total of 17 L of bacterial cultures were cultivated and centrifuged at 10,000 g for 10 min to remove cells. The culture supernatants were extracted twice with equal volumes of ethyl acetate. The resultant organic layer was concentrated *in vacuo* to generate a total crude extract of 1.5 g.

The crude extract was adsorbed onto C_{18} resin (BAKERBONDTM Octadecyl (C_{18}) 40 µm Prep LC Packing; J.T. Baker, Phillipsburg, NJ) and separated by reversed-phase flash chromatography. A linear gradient of 10 – 100% MeOH in H₂O, each solvent modified with 0.1% formic acid (FA), afforded five fractions. Fraction 4 (135.6 mg) was further purified by reversed-phase C_{18} semi-preparative HPLC (25 min linear gradient from 40 – 78% MeOH in H₂O (0.1% FA) to yield 5 pure compounds: **1** (28.3 mg, t_R = 17.9 min), **2** (10.1 mg, t_R = 16.9 min), **3** (7.8 mg, t_R = 13.8 min), **4** (2.3 mg, t_R = 16.0 min), and **5** (2.2 mg, t_R = 15.1 min).

Thalassotalic acid A (1): off-white amorphous powder; UV (MeOH) λ_{max} (log ε) 201 (4.16) 298 (3.99) nm; IR (ZnSe) v_{max} 3308, 3067 (br), 2917, 2850, 1688, 1656, 1601, 1584, 1271, 1167 cm^{-1; 1}H and ¹³C NMR data, see Table 1; HRESIMS *m/z* 332.1866 [M-H]⁻, (calcd for C₁₉H₂₆NO₄, 332.1867).

Thalassotalic acid B (2): brown amorphous powder; UV (MeOH) λ_{max} (log e) 201 (4.12) 299 (4.03) nm; IR (ZnSe) v_{max} 3326, 3118 (br), 2925, 2853, 1688, 1656, 1601, 1584, 1270, 1168 cm^{-1; 1}H and ¹³C NMR data, see Table 1; HRESIMS *m/z* 332.1872 [M-H]⁻, (calcd for C₁₉H₂₆NO₄, 332.1867).

Thalassotalic acid C (3): brown amorphous powder; UV (MeOH) λ_{max} (log e) 202 (4.27) 299 (4.37) nm; IR (ZnSe) ν_{max} 3317, 3067 (br), 2922, 2852, 1692, 1655, 1601, 1584, 1270, 1168 cm^{-1; 1}H and ¹³C NMR data, see Table 1; HRESIMS *m/z* 318.1699 [M-H]⁻, (calcd for C₁₈H₂₄NO₄, 318.1711).

Thalassotalamide A (4): brown amorphous powder; ¹H and ¹³C NMR data, see Table 1; HRESIMS m/z 331.2021 [M-H]⁻, (calcd for C₁₉H₂₇N₂O₃, 331.2027).

Thalassotalamide B (5): brown amorphous powder; UV (MeOH) λ_{max} (log ε) 200 (4.26) 210 (4.19) 272 (3.59) nm; IR (ZnSe) v_{max} 3726, 3624, 3321, 2925, 2854, 1665, 1578, 1137 cm^{-1; 1}H and ¹³C NMR data, see Table 1; HRESIMS *m/z* 331.2015 [M-H]⁻, (calcd for C₁₉H₂₇N₂O₃, 331.2027).

Tyrosinase Inhibition Assay

Inhibitory effects of compounds 1–5 on mushroom tyrosinase were evaluated spectrophotometrically using L-tyrosine as a substrate according to our previously published method with minor modification.²⁵ Tyrosinase inhibition assays were performed in 96-well microplate format using a SpectraMax M2 microplate reader (Molecular Devices, CA). Briefly, compounds were dissolved at various concentrations in 10% MeOH in phosphate buffer solution (PBS; 0.1 M, pH 6.8). Each well contained 40 μ L of sample with 80 μ L of PBS, 40 μ L of tyrosinase (100 units/mL), and 40 μ L L-tyrosine (2.5 mM). The mixture was incubated for 30 min at 37 °C, and absorbance was measured at 490 nm. Each sample was accompanied by a blank containing all components except L-tyrosine. Arbutin and kojic acid were used as positive controls. The results were compared with a control consisting of 10% MeOH in place of the test compounds. Experiments were completed in triplicate for each compound. The percentage of tyrosinase inhibition was calculated as: [(Acontrol – Asample)/ Acontrol] × 100.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

Instruments used for mass spectrometric and other analyses were supported through an Institutional Development Award (IDeA) from the National Institute of General Medical Sciences of the National Institute of Health (grant number 2 P20 GM103430). NMR data were acquired on instruments supported in part by the National Science Foundation EPSCoR Cooperative Agreement #EPS-1004057. J.C. was supported in part by the China Scholarship Council (File no. 201408330156).

References

 Blunt JW, Copp BR, Keyzers RA, Munro MH, Prinsep MR. Nat Prod Rep. 2015; 32:116–211. [PubMed: 25620233]

- Blunt JW, Copp BR, Keyzers RA, Munro MH, Prinsep MR. Nat Prod Rep. 2014; 31:160–258. [PubMed: 24389707]
- 3. Blunt JW, Copp BR, Keyzers RA, Munro MH, Prinsep MR. Nat Prod Rep. 2013; 30:237–323. [PubMed: 23263727]
- Machado H, Sonnenschein EC, Melchiorsen J, Gram L. BMC genomics. 2015; 16:158. [PubMed: 25879706]
- 5. Still PC, Johnson TA, Theodore CM, Loveridge ST, Crews P. J Nat Prod. 2014; 77:690–702. [PubMed: 24571234]
- 6. Tsukimoto M, Nagaoka M, Shishido Y, Fujimoto J, Nishisaka F, Matsumoto S, Harunari E, Imada C, Matsuzaki T. J Nat Prod. 2011; 74:2329–2331. [PubMed: 22035372]
- Xu Y, Kersten RD, Nam SJ, Lu L, Al-Suwailem AM, Zheng H, Fenical W, Dorrestein PC, Moore BS, Qian PY. J Am Chem Soc. 2012; 134:8625–8632. [PubMed: 22458477]
- Sudek S, Lopanik NB, Waggoner LE, Hildebrand M, Anderson C, Liu H, Patel A, Sherman DH, Haygood MG. J Nat Prod. 2007; 70:67–74. [PubMed: 17253852]
- 9. Mansson M, Gram L, Larsen TO. Mar Drugs. 2011; 9:1440-68. [PubMed: 22131950]
- 10. Isnansetyo A, Kamei Y. J Ind Microbiol Biotechnol. 2009; 36:1239-48. [PubMed: 19582493]
- 11. Zhang Y, Tang K, Shi X, Zhang XH. Int J Syst Evol Micr. 2014; 64:1223–1228.
- Stelling SC, Techtmann SM, Utturkar SM, Alshibli NK, Brown SD, Hazen TC. Genome Announc. 2014; 2:e01231–14.
- 13. Park S, Jung YT, Kang CH, Park JM, Yoon JH. Int J Syst Evol Micr. 2014; 64:3676–3682.
- 14. Hou T-T, Liu Y, Zhong Z-P, Liu H-C, Liu Z-P. Int J Syst Evol Micr. 2015
- 15. Thompson F, Barash Y, Sawabe T, Sharon G, Swings J, Rosenberg E. Int J Syst Evol Micr. 2006; 56:365–368.
- Ohta Y, Hatada Y, Miyazaki M, Nogi Y, Ito S, Horikoshi K. Curr Microbiol. 2005; 50:212–216. [PubMed: 15902469]
- 17. Hearing VJ, Tsukamoto K. FASEB J. 1991; 5:2902-2909. [PubMed: 1752358]
- Ando H, Kondoh H, Ichihashi M, Hearing VJ. J Invest Dermatol. 2007; 127:751–761. [PubMed: 17218941]
- 19. Parvez S, Kang M, Chung HS, Bae H. Phytother Res. 2007; 21:805–816. [PubMed: 17605157]
- 20. Torres M, Romero M, Prado S, Dubert J, Tahrioui A, Otero A, Llamas I. Microbiol Res. 2013;
 - 168:547–554. [PubMed: 23743010]
- 21. Guglielmini G. J Appl Cosmetol. 2006; 24:55–61.
- 22. Guglielmini G. Cosmet Toiletries. 2004; 119:61–65.
- 23. Kafara M, Arct J, Dzierzgowski S. J Appl Cosmetol. 2009; 27:1–12.
- 24. Alvaro, M., Cozzi, R., Genovese, A., Sedghi Zadeh, H., Patent, I. WO. 2013/179098 A1. 2013.
- Niesen DB, Ma H, Yuan T, Bach A 2nd, Henry GE, Seeram NP. Nat Prod Commun. 2015; 10:491– 493. [PubMed: 25924536]
- 26. Chang TS. Int J Mol Sci. 2009; 10:2440-2475. [PubMed: 19582213]
- Wang YH, Avonto C, Avula B, Wang M, Rua D, Khan IA. J AOAC Int. 2015; 98:5–12. [PubMed: 25857872]
- 28. Brady SF, Clardy J. Org Lett. 2005; 7:3613–3616. [PubMed: 16092832]
- 29. Brady SF, Chao CJ, Clardy J. Appl Environ Microb. 2004; 70:6865–6870.
- 30. Van Wagoner RM, Clardy J. Structure. 2006; 14:1425-1435. [PubMed: 16962973]

Author Manuscript



Figure 1. Key COSY, HMBC, and NOESY correlations for thalassotalic acid A (1).

Table 1

NMR spectroscopic data (500 MHz, DMSO- d_6) for thalassotalic acids A–C (1–3) and thalassotalamides A (4) and B (5).

	Thalassotalic aci	id A (1)	Thalassotal	ic acid B (2)	Thalassotalic aci	d C (3)	Thalassotalamide	e A (4)	Thalassotals	unide B (5)
position	b c	$\delta_{\rm H} (J \text{ in Hz})$	δ_{C}	$\delta_{ m H}(J{ m in}{ m Hz})$	ઈ C	$\delta_{ m H}~(J~{ m in}~{ m Hz})$	δ _C	$\delta_{\rm H} (J \text{ in Hz})$	S C	$\delta_{\rm H} (J \text{ in Hz})$
-	166.7, C		166.7, C		166.8, C		167.2, C		167.2, C	
1-OH/NH ₂		12.35, br s		12.23, br s		12.26, br s		7.28/7.05, br s		7.28/7.05, br s
2	124.2, C		124.3, C		124.5, C		127.0, C		127.0, C	
2-NH		9.23, s		9.22, s		9.20, s		9.18, s		9.19, s
3	132.2, CH	7.22, s	132.0, CH	7.22, s	131.7, ^a CH	7.21, s	128.6, CH	7.06, s	128.6, CH	7.06, s
4	124.7, C		124.7, C		124.8, C		125.0, C		125.0, C	
5	131.7, CH	7.51, d (8.7)	131.7, CH	7.51, d (8.6)	131.7, ^a CH	7.50, d (8.5)	131.2, CH	7.42, d (8.7)	131.2, CH	7.42, d (8.5)
9	115.3, CH	6.80, d (8.7)	115.3, CH	6.80, d (8.5)	115.3, CH	6.79, d (8.5)	115.2, CH	6.78, d (8.7)	115.2, CH	6.78, d (8.5)
7	158.5, C		158.5, C		158.5, C		158.0, C		158.1, C	
HO-7		9.94, br s		10.02, br s		10.03, br s		10.00, br s		10.03, br s
1′	172.0, C		171.9, C		171.8, C		172.0, C		172.0, C	
2,	35.1, CH ₂	2.28, t (7.2)	$35.2, CH_2$	2.28, t (7.4)	35.2, CH ₂	2.28, t (7.2)	35.2, CH ₂	2.32, t (7.5)	35.2, CH ₂	2.32, t (7.5)
3,	$25.0, CH_2$	1.59, m	$25.0, CH_2$	1.59, m	25.0, CH ₂	1.59, m	24.7, CH ₂	1.58, m	24.7, CH ₂	1.58, m
,4	28.6–29.0, CH ₂	1.31, m ^a	28.7, CH ₂	1.34, m ^{<i>a</i>}	28.6–28.8, CH ₂	1.31, m ^a	28.7–28.9, CH ₂	1.31, m ^{<i>a</i>}	28.8, CH ₂	1.32, m ^{<i>a</i>}
5,	28.6–29.0, CH ₂	1.31, m ^a	29.1, CH ₂	1.30, m ^{<i>a</i>}	28.6–28.8, CH ₂	1.31, m ^a	28.7–28.9, CH ₂	1.31, m ^{<i>a</i>}	29.1, CH ₂	1.28, m ^{<i>a</i>}
6′	28.6–29.0, CH ₂	1.31, m ^a	26.7, CH ₂	1.30, m ^{<i>a</i>}	28.6–28.8, CH ₂	1.31, m ^a	28.7–28.9, CH ₂	1.31, m ^{<i>a</i>}	26.7, CH ₂	1.30, m ^{<i>a</i>}
7′	28.6–29.0, CH ₂	1.31, m ^a	38.4, CH ₂	1.19, m	31.2, CH ₂	1.30, m ^{<i>a</i>}	28.7–28.9, CH ₂	1.31, m ^{<i>a</i>}	38.4, CH ₂	1.19, m
×,	31.3, CH ₂	1.29, m ^{<i>a</i>}	27.4, CH ₂	1.55, m	22.1, CH ₂	1.32, m ^{<i>a</i>}	31.3, CH ₂	1.29, m ^{<i>a</i>}	27.4, CH ₂	1.55, m
9,	22.1, CH ₂	1.30, m ^{<i>a</i>}	22.5, CH ₃	0.90, d (6.6)	13.9, CH ₃	0.90, t (7.0)	22.1, CH ₂	1.30, m ^{<i>a</i>}	22.5, CH ₃	0.89, d (6.5)
10'	13.9, CH ₃	0.90, t (7.0)					13.9, CH ₃	0.90, t (7.0)		
^a Overlapped si	ignal									

Table 2

Tyrosinase inhibitory activity of compounds 1–5.

Compound	IC ₅₀ (µM)
1	130 ± 10
2	470 ± 10
3	280 ± 10
4	> 1000
5	> 1000
6 ^a	100 ± 10
7 ^{<i>a</i>}	46 ± 2

^aPositive controls