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In Silico Prediction of Anti-Inflammatory Potential of Isolated Compounds from Enhalus acoroides

Thuy X. Ho¹, Tuan N. Nguyen², Hang T.T. Tran¹, Tram T.T. Nguyen¹, Idania Rodeiro Guerrae³, Phuong L. Doan^{4,5}, Thang D. Tran²*

¹Department of Chemistry, Vinh University, Vinh City, Nghean, Vietnam

²Institute of Biotechnology and Food Technology, Industrial University of Ho Chi Minh City, Ho Chi Minh City, Vietnam

³Institute of Marine Sciences, Ministry of Science, Technology and Environmental, Havana, Cuba

⁴Institute of Natural Products Chemistry, Vietnam Academy of Science and Technology, 18 Hoang Quoc Viet, Cau Giay, Hanoi, Vietnam

⁵Graduate University of Science and Technology, Vietnam Academy of Science and Technology, 18 Hoang Quoc Viet, Cau Giay, Hanoi, Vietnam

ARTICLE INFO	ABSTRACT
Article history:	Enhalus acoroides is a marine flowering plant with numerous biological activities. The current
Received 10 November 2024	study aim to isolate, characterize the chemical constituents from Enhalus acoroides whole plant,
Revised 17 November 2024	and to evaluate their anti-inflammatory activity in silico. The dried powdered E. acoroides whole
Accepted 15 January 2025	plant was subjected to extraction, portioning, and chromatographic separation for the isolation of
Published online 01 March 2025	its phytochemical constituents. The structures of the isolated compounds were elucidated by a
Copyright: © 2025 Ho <i>et al.</i> This is an open-access article distributed under the terms of the <u>Creative</u> <u>Commons</u> Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.	combination of Nuclear Magnetic Resonance (NMR) spectroscopy, Mas Spectrometry (MS), and comparison of spectral data with literature. The isolated compounds were investigated for their potential anti-inflammatory activity <i>in silico</i> via molecular docking of the compounds with inducible nitric oxide synthase (iNOS). Phytochemical investigation of the whole plant of <i>E.</i> <i>acoroides</i> led to the isolation of ten (10) compounds. Their chemical structures were determined as methyl pheophorbide a (1), (+)-catechin-4'-O- β -D-glucoside (2), quercetin (3), rutin (4), apigenin-7-O- α -L-rhamnopyranosyl-(1 \rightarrow 2)- β -D-glucopyranoside (5), adenosine (6), uracil (7), D-mannitol (8), 6β -hydroxystigmast-4-en-3-one (9), and daucosterol (10) by ESI-MS, NMR spectral analysis, and comparison with published data. Molecular docking simulations of the compounds revealed their potential iNOS inhibitory activity, with compounds 1, 2, 3, 4, 5, 9, and 10 showing stronger binding affinities than the reference inhibitor, ethyl 4-[(4-methylpyridin-2- yl)amino]piperidine-1-carboxylate (AR-C95791). These findings suggest that the compounds, particularly compound 10, hold significant promise as iNOS inhibitors, potentially serving as candidates for anti-inflammatory drug development.

Keywords: Enhalus acoroides, Chemical constituents, Anti-inflammatory, Binding affinity.

Introduction

Enhalus acoroides (L.f.) Royle, belonging to the Hydrocharitaceae family, is a marine flowering plant that grows in the shallow coastal waters of tropical and temperate regions.¹ In Vietnam, *E. acoroides* is commonly found in the Tam Giang-Cau Hai lagoon area (Thua Thien-Hue province) and Cam Lam Beach (Khanh Hoa province). Phytochemical screening of *E. acoroides* has revealed the presence of flavonoids, steroids, monoterpenoids, diterpenoids, and aliphatic acids.²⁻⁴ Pharmacological investigations have demonstrated that *E. acoroides* exhibits various biological activities such as antioxidant, antifeedant, antibacterial, larvicidal, antitumour, and cytotoxic effects.¹⁻⁵ However, the anti-inflammatory activity of *E. acoroides* has not been studied.

*Corresponding author. Email: <u>thangtd@iuh.edu.vn</u> Tel: +84-91-3049689.

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In a targeted bioprospecting effort to discover new anti-inflammatory agents from marine flowering plants, the present study reports the isolation and characterization of ten compounds from *E. acoroides* collected in Vietnam. An *in silico* study was conducted to first evaluate the potential binding affinity of the isolated compounds for the anti-inflammatory inducible nitric oxide synthase (iNOS) target protein. The results from the *in silico* study is aimed at predicting potential active compounds in *Enhalus acoroides* that could further be investigated *in vitro* and *in vivo*.

Materials and Methods

General experimental procedures

The NMR data were recorded using Bruker AV-III 500 NMR spectrometer with TMS as the internal standard and chemical shifts were reported in δ values (ppm). The electrospray ionization mass spectrometry (ESI-MS) spectra were measured on an Agilent LC/MS-6420. The preparative HPLC was run on an Agilent 218 Purification System using the ZORBAX SB-C18 columns (100 mm × 21.2 mm × 5µm). Silica gel 230–400 mesh (0.040–0.063 mm, Merck) and YMC RP-18resins (30–50 µm, Fujisilisa Chemical Ltd.) were used for the column chromatography (CC). The thin-layer chromatography (TLC) was done on pre-coated silica gel 60-F254 plates. Spots were visualized by spraying with 10% sulfuric acid reagent, and heating. All chemicals and reagents used in this study were of analytical grade.

Plant materials

The whole plant of *E. acoroides* was collected from the Van Phong-Khanh Hoa province (12.1960803 N, 108.9950386 E), Vietnam in March 2023. The plant material was identified and authenticated by Assoc. Prof. Dr. Dam Duc Tien, a plant taxonomist in the Department

of Chemistry, Vinh University, Vietnam. A herbarium specimen with voucher No. XuanThuy-032021 was deposited at the Laboratory of the department.

Extraction and isolation

The air-dried whole plant of E. acoroides (14 kg) were cut into small pieces and extracted with methanol (50 L \times 3 times) by maceration at room temperature. The solvent was removed by a vacuum evaporator to yield the crude methanol extract (685 g), which was further suspended in water and successively partitioned with n-hexane (8 L × 3 times), ethyl acetate (8 L \times 3 times) and n-butanol (8 L \times 3 times) to give n-hexane extract (EAH 157.5 g), ethyl acetate extract (EAE 238.9 g), and n-butanol extract (EAB 64.6 g), respectively. The n-hexane extract was subjected to silica gel chromatography column (CC), eluted with hexane/acetone (100:1 to 1:1, v/v) to yield seven fractions (Frs. EAH1 - EAH7). Fraction EAH3 (6.1 g) was subjected to silica gel CC eluted with hexane/ethyl acetate (50:1 to 2:1, v/v) to afford compound 9 (22.4 mg). The ethyl acetate extract was separated over a silica gel CC, eluted with chloroform/methanol (100:0 to 3:1, v/v) to produce eight fractions (Frs. EAE1-EAE8). Fraction EAE2 (16.4 g) was applied to silica gel CC, eluted with n-hexane-acetone (10:1 to 1:1, v/v) to afford compound 1 (44.7 mg). Fraction EAE3 (11.5 g) was subjected to a silica gel CC, eluted with dichloromethane/methanol (25:1 to 0:1, v/v) to yield four subfractions (Frs. EAE3.1-EAE3.4). Subfraction EAE3.3 (2.9 g) was further purified using preparative HPLC (60% methanol in water) to obtain compound 3 (20.1 mg). The n-butanol extract (EAB) was also subjected to silica gel CC eluted with gradient chloroform/methanol/water (50:1:0 to 1:1:1, v/v) to obtain nine fractions (Frs. EAB1 - EAB9). Further purification of fraction EAB2 (1.2 g) was separated on silica gel CC, eluted with chloroform/methanol (90:10 to 25:75 v/v) to give compound 4 (45.9 mg). Fraction EAB3 (45.1 mg) was purified using preparative HPLC (70% acetonitrile in water) to yield compound 5 (10.4 mg). Subfraction EAB4 (1.7 g) was separated by CC with RP-C18 eluted with methanol:water (3:1), and further purified by sephadex LH-20 CC to give compound 6. Compound 2 (42.3 mg) was isolated from subfraction EAB5 (8.1 g) by silica gel CC eluted with chloroform/methanol (7:1, 4:1, 1:1 v/v in a stepwise gradient). Fraction EAB6 (2.1 g) was purified using preparative HPLC (50% acetonitrile in water) to yield compound 7 (9.6 mg) and compound 8 (11.2 mg). The EAB8 fraction (4.2 g) was separated by CC eluted with chloroform:methanol (9:1) to obtain compound 10 (53.7 mg).

Molecular docking

The structure of inducible nitric oxide synthase in complex with AR-C95791 was obtained from the RCSB Protein Data Bank (https://www.rcsb.org/structure/3E7G) (PDB ID: 3E7G).⁶ All ions, water molecules, and co-crystallized ligands were removed, and hydrogen atoms along with Kollman charges were added using AutoDockTools v1.5.6. The 3D structures of the isolated compounds were drawn using Marvin JS software, then their energy was optimized using the UFF force field in the OpenBabel program package.7, Subsequently, the ligand and protein files were converted to PDBQT format, ready for molecular docking simulations using AutoDock Vina v1.2.3.9,10 The box coordinate parameters were set as follows: center coordinates were defined as x = 55 Å, y = 21.8 Å, z = 78.7 Å, box size X x Y x Z = 24 x 24 x 24, spacing = 1, and other parameters were set to default. The exhaustiveness value was adjusted to 400 as reported previously.^{11,12} The re-docking process was performed with the cocrystallized ligand AR-C95791 (Ethyl 4-[(4-methylpyridin-2yl)amino]piperidine-1-carboxylate) to verify the parameters for a reliable binding mode prediction. After obtaining the docking results of the isolated compounds with the iNOS protein target, Discovery Studio Visualizer was used to analyze and visualize their interactions.

Results and Discussion

Isolation and structure elucidation

The column chromatographic separation, and preparative HPLC of the crude methanol extract of *Enhalus acoroides* whole plant led to the isolation of ten compounds. These compounds included methyl pheophorbide a (1), (+)-catechin-4'- $O-\beta$ -D-glucoside (2), quercetin (3),

rutin (4), apigenin-7-O- α -L-rhamnopyranosyl-(1 \rightarrow 2)- β -D-glucopyranoside (5), adenosine (6), uracil (7), D-mannitol (8), 6β - hydroxystigmast-4-en-3-one (9) and daucosterol (10).

Compound 2 was obtained as a white amorphous powder. The ESI-MS spectrum of 2 exhibited a molecular ion peak at m/z 453 [M+H]⁺, corresponding to the molecular formula of C₂₁H₂₄O₁₁. The ¹H-NMR spectra of compound 2 exhibited signals of five aromatic protons at $\delta_{\rm H}$ 6.90 – 5.80 ppm and an anomeric proton signal at $\delta_{\rm H}$ 4.77 (1H, d, J = 6.9 Hz, H-1''). The ¹³C-NMR spectrum of compound 2 exhibited 21 carbon signals, including 15 of a flavan-3-ol and 6 of a sugar, suggesting a flavonoid glycoside. The NMR data of compound 2 were compared with the literature data reported for (+)-catechin-4'-*O*- β -D-glucoside, ¹³ and were found to match.

Compound 3 was obtained as a yellow amorphous powder. The ESI-MS spectrum of 3 exhibited a molecular ion peak at m/z 301 [M-H]⁻, corresponding to the molecular formula of $C_{15}H_{10}O_7$. The ¹H-NMR spectra of 3 exhibited five aromatic protons at δ_H 7.76 (1H, s, H-2'), 7.66 (1H, d, J = 8.0 Hz, H-6'), 6.91 (1H, d, J = 8.0 Hz, H-5'), 6.41 (1H, s, H-8), and 6.21 (1H, s, H-6). The ¹³C-NMR spectra of 3 showed fifteen carbon signals including a carbonyl group at δ_C 177.4 (C-4), seven oxygenated aromatic carbons at δ_C 165.6 (C-7), 162.5 (C-5), 158.3 (C-9), 148.0 (C-2/C-4'), 146.2 (C-3'), 137.2 (C-3), two quaternary aromatic carbons at δ_C 124.2 (C-1') and 104.5 (C-10), five sp² methine carbons at δ_C 121.7 (C-6'), 116.2 (C-5'), 116.0 (C-2'), 99.3 (C-6), and 94.4 (C-8) were confirmed a flavonoid bearing five hydroxyl groups. Based on NMR, MS data, and published data,¹⁴ compound 3 was characterized as quercetin.

Compound 4 was obtained as a yellow amorphous powder. Its ESI-MS spectrum of 4 showed a molecular ion peak at m/z 611 $[M+H]^+$, corresponding to the molecular formula of $C_{27}H_{30}O_{16}$. The ¹H-NMR spectra of 4 exhibited signals of two aromatic systems, characteristics of the quercetin compound. The ¹³C-NMR spectra of 4 revealed 27 carbon signals, including 15 carbons of a flavonol skeleton and 12 carbon of two sugar units, suggesting a flavonoid glycoside. Based on the MS, NMR analyses, and published data,¹⁵ compound 4 was elucidated as rutin.

The other compounds were elucidated as methyl pheophorbide a (1),¹⁶ apigenin-7-O- α -L-rhamnopyranosyl-(1 \rightarrow 2)- β -D-glucopyranoside (5),¹⁷ adenosine (6),¹⁸ uracil (7),¹⁹ D-mannitol (8),²⁰ 6 β -hydroxystigmast-4-en-3-one (9),²¹ and daucosterol (10).²² Their MS and NMR data were compared to those of the literature and found to match. To the best of our knowledge, this is the first report on the isolation and structure elucidation of chemical constituents from *Enhalus acoroides*. The structures of these components are shown in Figure 1.

Characterization of the isolated compounds

Methyl pheophorbide a (1): white amorphous powder; ESI-MS: m/z 607 [M+H]⁺; ¹H-NMR (500 MHz, CDCl₃) δ_H (ppm): 9.46 (1H, s, H-10), 9.30 (1H, s, H-5), 8.54 (1H, s, H-20), 7.93 (1H, dd, J = 18.0, 11.5 Hz, H-3'), 6.25 (1H, s, H-13''), 6.25 (1H, d, J = 17.5 Hz, H-3''E), 6.15 (1H, dd, J = 12.0, 1.0 Hz, H-3"Z), 4.45 (1H, dt, J = 7.0, 1.5 Hz, H-18), 4.20 (1H, d, J = 10.0 Hz, H-17), 3.88 (3H, s, H-13'''), 3.66 (3H, s, H-12'), 3.62 (2H, q, J = 7.5 Hz, H-8'), 3.57 (3H, s, H-17''''), 3.37 (3H, s, H-2'), 3.17 (3H, s, H-7'), 2.63 (1H, m, H-17'a), 2.51 (1H, m, H-17"a), 2.32 (1H, m, H-17'b), 2.25 (1H, m, H-17'b), 1.81 (3H, d, *J* = 7.5 Hz, H-18'), 1.66 (3H, t, J = 7.5 Hz, H-8''); ¹³C-NMR (125 MHz, CDCl₃) $\delta_{\rm C}$ (ppm): 189.6 (C-13'), 173.4 (C-17'''), 172.2 (C-19), 169.6 (C-13'''), 161.2 (C-16), 155.6 (C-6), 151.0 (C-9), 149.7 (C-14), 145.2 (C-8), 142.1 (C-1), 137.9 (C-11), 136.5 (C-3), 136.2 (C-4), 136.1 (C-7), 131.9 (C-2), 129.1 (C-12), 129.0 (C-13/C-3'), 122.8 (C-3''), 105.2 (C-15), 104.4 (C-10), 97.5 (C-5), 93.1 (C-20), 64.7 (C-13"), 52.8 (C-13""), 51.7 (C-17""), 51.1 (C-17), 50.1 (C-18), 31.1 (C-17"), 29.9 (C-17"), 23.1 (C-18'), 19.4 (C-8'), 17.4 (C-8''), 12.1 (C-2'/C-12'), 11.2 (C-7').



Figure 1: Structures of isolated compounds (1 - 10) from *Enhalus acoroides* whole plant

(+)-*Catechin-4* '-*O-β* -*D*-glucoside (2): white amorphous powder; ESI-MS: m/z 453 [M+H]⁺; ¹H-NMR (500 MHz, CD₃OD) $\delta_{\rm H}$ (ppm): 6.86 (1H, d, J = 2.0 Hz, H-2'), 6.79 (1H, m, H-5'), 6.75 (1H, dd, J = 8.0, 2.0 Hz, H-6'), 5.95 (1H, d, J = 2.0 Hz, H-6), 5.88 (1H, d, J = 2.5 Hz, H-8), 4.77 (1H, d, J = 6.9 Hz, H-1''), 4.59 (1H, d, J = 7.5 Hz, H-2), 4.00 (1H, m, H-3), 3.80 – 3.50 (4H, m, H-2''/H-3''/H-4''/H-5''), 3.61 (1H, dd, J = 11.0, 5.0 Hz, H-6''a), 3.54 (1H, dd, J = 11.0, 6.0 Hz, H-6''b), 2.87 (1H, dd, J = 16.5, 5.5 Hz, H-4a), 2.53 (1H, dd, J = 16.0, 8.0 Hz, H-4b); ¹³C-NMR (125 MHz, CD₃OD) $\delta_{\rm C}$ (ppm): 157.8 (C-7), 157.6 (C-5), 156.9 (C-9), 146.2 (C-3'/C-4'), 132.2 (C-1'), 120.0 (C-6'), 116.1 (C-5'), 115.3 (C-2'), 100.9 (C-10/C-1''), 96.3 (C-6), 95.5 (C-8), 82.8 (C-2), 79.9 (C-3''), 75.6 (C-2''), 73.9 (C-5''), 71.3 (C-4''), 68.8 (C-3), 64.4 (C-6''), 28.5 (C-4).

Quercetin (3): yellow amorphous powder; ESI-MS: m/z 301 [M-H]⁻; ¹H-NMR (500 MHz, CD₃OD) $\delta_{\rm H}$ (ppm): 7.76 (1H, s, H-2'), 7.66 (1H, d, *J* = 8.0 Hz, H-6'), 6.91 (1H, d, *J* = 8.0 Hz, H-5'), 6.41 (1H, s, H-8), 6.21 (1H, s, H-6); ¹³C-NMR (125 MHz, CD₃OD) $\delta_{\rm C}$ (ppm): 177.4 (C-4), 165.6 (C-7), 162.5 (C-5), 158.3 (C-9), 148.0 (C-2/C-4'), 146.2 (C-3'), 137.2 (C-3), 124.2 (C-1'), 121.7 (C-6'), 116.2 (C-5'), 116.0 (C-2'), 104.5 (C-10), 99.3 (C-6), 94.4 (C-8).

Rutin (4): yellow amorphous powder; ESI-MS: m/z 611 [M+H]⁺; ¹H-NMR (500 MHz, DMSO-d₆) $\delta_{\rm H}$ (ppm): 12.59 (1H, s, 5-OH), 7.54 (1H, dd, J = 8.4, 2.0 Hz, H-6'), 7.53 (1H, d, J = 2.0 Hz, H-2'), 6.83 (1H, d, J = 8.0 Hz, H-5'), 6.38 (1H, d, J = 2.0 Hz, H-8), 6.19 (1H, d, J = 2.0 Hz, H-6), 5.15 (1H, d, J = 7.2 Hz, H-1''), 4.56 (1H, d, J = 1.6 Hz, H-1'''), 3.70 (1H, d, J = 10.5 Hz, H-6''a), 3.30 (1H, m, H-6''b), 3.29 (1H, m, H-3'''), 3.28 (1H, m, H-2'''), 3.25 (1H, m, H-5'''), 3.24 (1H, m, H-5''), 3.22 (1H, m, H-2''), 3.21 (1H, m, H-3'''), 3.07 (1H, m, H-4'''), 3.05 (1H, m, H-4'''), 0.99 (3H, d, J = 6.2 Hz, H-6''); ¹³C-NMR (125 MHz, DMSO-d₆) $\delta_{\rm C}$ (ppm): 177.4 (C-4), 164.1 (C-7), 161.2 (C-5), 156.6 (C-2), 156.4 (C-9), 148.4 (C-4'), 144.7 (C-3'), 135.3 (C-3), 121.6 (C-1'), 121.2 (C-6'), 116.3 (C-2'), 115.2 (C-5'), 103.9 (C-10), 101.2 (C-1''), 100.7 (C-1'''), 98.7 (C-6), 93.6 (C-8), 76.4 (C-3''), 76.0 (C-5''), 74.1 (C-2''), 71.8 (C-4''), 70.6 (C-3'''), 70.4 (C-2'''), 70.0 (C-4''), 68.5 (C-5'''), 67.0 (C-6'''), 17.7 (C-6''').

Apigenin-7-O-α-L-rhamnopyranosyl-($1 \rightarrow 2$)- β -*D-glucopyranoside* (5): yellow amorphous powder; ESI-MS: m/z 579 [M+H]⁺; ¹H-NMR (500 MHz, CD₃OD) $\delta_{\rm H}$ (ppm): 7.89 (2H, d, J = 8.5 Hz, H-2'/H-6'), 6.95 (2H, d, J = 8.5 Hz, H-3'/H-5'), 6.80 (1H, s, H-8), 6.66 (1H, s, H-3), 6.47 (1H, s, H-6), 5.31 (1H, s, H-1'''), 5.21 (1H, d, J = 8.0 Hz, H-1''), 1.33 (3H, d, J = 6.5 Hz, H-6'''); ¹³C-NMR (125 MHz, CD₃OD) $\delta_{\rm C}$ (ppm): 184.1 (C-4), 166.8 (C-2), 164.4 (C-5/C-7), 162.9 (C-4'), 159.0 (C-9), 129.7 (C-2'/C-6'), 123.1 (C-1'), 117.1 (C-3'/C-5'), 107.1 (C-10), 104.2 (C-3), 102.5 (C-1'''), 101.0 (C-6), 99.8 (C-1''), 95.9 (C-8), 79.0 (C-2''), 78.4 (C-3''), 78.3 (C-5''), 75.9 (C-4''), 74.0 (C-4''), 72.2 (C-3'''), 70.0 (C-5'''), 71.4 (C-2'''), 62.4 (C-6''), 18.3 (C-6''').

Adenosine (6): white amorphous powder; ESI-MS: m/z 266 [M-H]⁻; ¹H-NMR (500 MHz, CDCl₃ & CD₃OD) $\delta_{\rm H}$ (ppm): 8.23 (1H, s, H-2), 8.05 (1H, s, H-8), 5.86 (1H, d, J = 7.2 Hz, H-1'), 4.79 (1H, dd, J = 7.2, 4.8 Hz, H-2'), 4.35 (1H, dd, J = 4.8, 2.4 Hz, H-3'), 4.27 (1H, q, J = 2.4Hz, H-4'), 3.97 (1H, dd, J = 12.4, 2.4 Hz, H-5'a), 3.77 (1H, dd, J = 12.4, 2.4 Hz, H-5'b); ¹³C-NMR (125 MHz, CDCl₃ & CD₃OD) $\delta_{\rm C}$ (ppm): 156.2 (C-6), 152.5 (C-2), 148.6 (C-4), 141.0 (C-8), 120.6 (C-5), 91.3 (C-1'), 87.6 (C-4'), 74.2 (C-2'), 71.9 (C-3'), 61.6 (C-5').

Uracil (7): white amorphous powder; ESI-MS: m/z 111 [M-H]⁻; ¹H-NMR (DMSO-d₆, 500 MHz) $\delta_{\rm H}$ (ppm): 10.98 (1H, s, 3-NH), 10.81 (1H, s, 1-NH), 7.37 (1H, d, J = 7.5 Hz, H-6), 5.44 (1H, d, J = 7.5 Hz, H-5).

D-mannitol (8): white amorphous powder; ESI-MS: m/z 183 [M+H]⁺; ¹H-NMR (500 MHz, DMSO-d₆) $\delta_{\rm H}$ (ppm): 3.61 (2H, m, H-1a/H-6a), 3.53 (2H, t, *J* = 7.5 Hz, H-3/H-4), 3.45 (2H, m, H-2/H-5), 3.37 (2H, dd, *J* = 11.5, 6.0 Hz, H-1b/H-6b); ¹³C-NMR (125 MHz, DMSO-d₆) $\delta_{\rm C}$ (ppm): 71.5 (C-2/C-5), 69.9 (C-3/C-4), 64.1 (C-1/C-6).

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1.20 – 1.12 (4H, m, H-12b/H-14/H-23), 1.03 (1H, m, H-22b), 1.02 (1H, m, H-17), 0.93 (2H, m, H-9/H-24), 0.92 (3H, d, J = 6.5 Hz, H-21), 0.85 (3H, t, J = 7.5 Hz, H-29), 0.84 (3H, d, 7.0 Hz, H-27), 0.82 (3H, d, J = 7.0 Hz, H-26), 0.74 (3H, s, H-18); ¹³C-NMR (125 MHz, CDCl₃) $\delta_{\rm C}$ (ppm): 200.4 (C-3), 168.5 (C-5), 126.3 (C-4), 73.3 (C-6), 56.1 (C-14), 55.9 (C-17), 53.7 (C-9), 45.9 (C-24), 42.6 (C-13), 39.7 (C-12), 38.6 (C-7), 38.0 (C-10), 37.1 (C-1), 36.1 (C-20), 34.3 (C-2), 34.0 (C-22), 29.8 (C-8), 29.2 (C-25), 28.2 (C-16), 26.2 (C-23), 24.2 (C-15), 23.1 (C-28), 21.0 (C-11), 19.8 (C-19), 19.5 (C-27), 19.1 (C-26), 18.8 (C-21), 12.0 (C-18/C-29).

Daucosterol (10): white amorphous powder; ESI-MS: m/z 414 [M-Glc]⁺; ¹H-NMR (500Hz, DMSO-d₆) $\delta_{\rm H}$ (ppm): 5.33 (1H, t, J = 4.5, 1.5 Hz, H-6), 4.86 (1H, d, J = 4.5 Hz, H-6'b), 4.84 (2H, t, J = 10.0, 4.5, Hz, H-3'/H-4'), 4.40 (1H, t, J = 11.5, 5.5, H-6a), 4.22 (1H, d, J = 7.5 Hz, H-1'), 3.66 - 2.88 (6H, m, H-2'/H-3'/H-4'/H-5'/H-6'a/H-6'b), 2.37 (1H, dd, J = 10.5, 2.5 Hz, H-4a), 2.13 (1H, s, H-4b), 1.97 (1H, m, H-12), 1.90 (1H, m, H-7), 1.82 (1H, m, H-2), 1.80 (2H, m, H-1/H-16), 1.61 (1H, m, H-25), 1.52 (1H, m, H-15), 1.50 (1H, m, H-7), 1.50 (1H, m, H-11), 1.47 (1H, m, H-2), 1.41 (1H, m, H-11), 1.39 (1H, m, H-8), 1.38 (1H, m, H-20), 1.30 (1H, m, H-22), 1.24 (1H, m, H-16), 1.22 (1H, m, H-28), 1.20 (1H, m, H-28), 1.18 (2H, m, H-23), 1,17 (1H, m, H-12), 1,10 (1H, m, H-17), 1.05 (1H, m, H-15), 0.99 (1H, m, H-22), 0.98 (1H, m, H-1), 0.97 (1H, m, H-14), 0.96 (3H, s, H-19), 0.93 (1H, m, H-24), 0.91 (1H, m, H-9), 0.90 (3H, d, J = 6.5 Hz, H-21), 0.82 (9H, m, H-26/H-27/H-29), 0.61 (3H, s, H-18); ¹³C-NMR (125 MHz, DMSO-d₆) δ_C (ppm): 140.4 (C-5), 121.1 (C-6), 100.8 (C-1'), 77.0 (C-3), 76.7 (C-5'), 76.6 (C-3'), 73.4 (C-2'), 70.1 (C-4'), 61.1 (C-6'), 56.2 (C-14), 55.4 (C-17), 49.6 (C-9), 45.1 (C-24), 41.8 (C-13), 40.1 (C-4), 40.0 (C-12), 38.2 (C-1), 36.8 (C-20), 36.2 (C-10), 35.4 (C-22), 33.3 (C-7), 31.4 (C-8), 31.3 (C-2), 29.2 (C-23), 28.7 (C-16), 27.7 (C-25), 25.5 (C-15), 23.8 (C-28), 22.6 (C-11), 20.6 (C-27), 19.7 (C-19), 19.0 (C-21), 18.9 (C-26), 11.7 (C-29), 11.7 (C-18).

Molecular docking data of isolated compounds

To investigate the inhibitory potential of compounds isolated from *Enhalus acoroides* whole plant against inducible nitric oxide synthases (iNOS), molecular docking was employed. iNOS plays an important

role in inflammation by producing nitric oxide (NO), an inflammatory mediator that helps protect the body against bacteria and harmful agents, but when overproduced, it can contribute to tissue damage and amplify the inflammatory response. By studying the interactions between these isolated compounds and iNOS, the drug design process can be optimized and deeper insights into the enzyme's role in inflammation pathology is gained.

The docking results are detailed in Tables 1 and 2, with the binding energies of compounds 1 - 10 ranging from -10.15 to -4.675 kcal/mol. Notably, seven of these compounds demonstrated stronger binding affinity than the reference inhibitor, AR-C95791 (-7.534 kcal/mol). Among the compounds, compound 10 emerged as the topmost candidate, showing the strongest binding to iNOS with an ΔG of -10.15 kcal/mol. This compound formed hydrogen bonds with Arg381 and Gln263, as well as pi-alkyl/alkyl interactions with Trp463, Val352, Arg199, Phe369, Cys200, Tyr489, Met355, Ala197, Leu209 and pisigma interactions with Trp194. Compound 5, with a binding energy of -10.15 kcal/mol, exhibited pi-sigma and pi-sulfur interactions with Met355, alongside pi-alkyl and alkyl interactions with Ala197, Cys200, and Val352. It also formed a hydrogen bond with Trp372 and pi-pi stacked interactions with Trp194. Compound 1 formed hydrogen bonds with Arg381 and Glu377, while also forming pi-anion interactions with Glu377, and pi-alkyl/alkyl interactions with Ile201, Val352, Pro350, Tyr347, and Glu377. Compound 3, which demonstrated a binding energy of -9.479 kcal/mol, formed hydrogen bonds with Tyr489 and engaged in pi-pi stacked and amide-pi stacked interactions with residues Trp194, Phe369, and Asn370. Meanwhile, compound 4 exhibited hydrogen bonds with Gln263, Arg109, Trp463, and Asp382, along with a pi-anion interaction with Glu377, and pi-alkyl/alkyl with Arg381, Pro350, contributing to a binding affinity of -8.696 kcal/mol. These results suggest that the compounds, particularly compound 10, hold significant promise as iNOS inhibitors, potentially offering a new avenue for the development of anti-inflammatory agents.

No.	Binding affinity (kcal/mol)	2D interaction	3D interaction
1	-9.554	ILE CYS ARG GY ASD ILE ASD GY ASD ILE ASD He A453 ASD ILE ASD He A453 ASD ILE ASD ASD ILE ASD	GLOSA PRO350 VAL352 ABG 381 TYR347

Table 1: Binding affinity, 2D- and 3D- interactions of the top compounds with the iNOS enzyme





No.	Compound	Binding affinity (kcal/mol)
1	Methyl pheophorbide a	-9.554
2	(+)-catechin-4'- O - β -D-glucoside	-8.195
3	quercetin	-9.479
4	rutin	-8.696
5	apigenin-7- O - α -L-rhamnopyranosyl-(1 \rightarrow 2)- β -D-glucopyranoside	-10.15
6	adenosine	-6.168
7	uracil	-4.675
8	D-mannitol	-4.846
9	6β -hydroxystigmast-4-en-3-one	-8.511
10	daucosterol	-10.12
11	Ethyl 4-[(4-methylpyridin-2-yl)amino]piperidine-1-carboxylate (AR- C95791)	-7.534

Table 2: The binding affinity of the isolated compounds 1-10 with the iNOS enzyme.

Conclusion

Phytochemical investigation of *E. acoroides* whole plant from Vietnam led to the isolation of ten (10) compounds including methyl pheophorbide a (1), (+)-catechin-4'-O- β -D-glucoside (2), quercetin (3), rutin (4), apigenin-7-O- α -L-rhamnopyranosyl-(1 \rightarrow 2)- β -Dglucopyranoside (5), adenosine (6), uracil (7), D-mannitol (8), 6β hydroxystigmast-4-en-3-one (9), and daucosterol (10). The structures of these compounds were determined by a combination of ESI-MS and NMR spectra and comparison with literature. Molecular docking was performed to evaluate the binding affinity of the isolated compounds with the iNOS enzyme. The findings suggest that compounds 1, 2, 3, 4, 5, 9, and 10 are promising iNOS enzyme inhibitors.

Conflict of Interest

The authors declare no conflict of interest.

Authors' Declaration

The authors hereby declare that the work presented in this article is original and that any liability for claims relating to the content of this article will be borne by them.

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