

Anti-HIV-1 Protease Triterpenoid Saponins from the Seeds of *Aesculus chinensis*

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Eight bioactive triterpenoid saponins (**1–8**) were isolated from the seeds of *Aesculus chinensis*, four of which are novel compounds. The major saponins were identified as escin Ia (**1**), Ib (**2**), isoescin Ia (**3**) and Ib (**4**), while the new compounds were identified as 22 α -tigloyl-28-acetylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1–2)] [β -D-glucopyranosyl (1–4)]- β -D-glucopyranosiduronic acid (escin IVc, **5**), 22 α -angeloyl-28-acetylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1–2)] [β -D-glucopyranosyl (1–4)]- β -D-glucopyranosiduronic acid (escin IVd, **6**), 28-tigloylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1–2)] [β -D-glucopyranosyl (1–4)]- β -D-glucopyranosiduronic acid (escin IVe, **7**), and 28-angeloylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1–2)] [β -D-glucopyranosyl (1–4)]- β -D-glucopyranosiduronic acid (escin IVf, **8**). The structures were determined by chemical and spectroscopic methods. All the above compounds were evaluated for their inhibitory activity against HIV-1 protease.

Aesculus chinensis Bge. (Hippocastanaceae) is a medicinal plant widely distributed in northwestern China. Its dried ripe seeds have been used as a carminative, stomachic, and analgesic for the treatment of distention and pain in chest and abdomen.¹ The saponin mixtures extracted from the seeds are known as escins, which have distinguished antiinflammatory, antiedema, capillary-protective, hypoglycemic and ethanol absorption inhibitory activities.^{2–4} In the course of our search for novel anti-HIV agents, escins were found to show moderate anti HIV-1 protease activity. Here we describe the isolation and structure elucidation of four novel triterpenoid saponins, escin IVc (**5**), IVd (**6**), IVe (**7**), and IVf (**8**). All of the isolated compounds were investigated for inhibitory activities against HIV-1 protease.

Results and Discussion

EtOH extracts of the seeds of Chinese horse chestnut were separated as described in the Experimental Section to yield compounds **1–8**. The four major active compounds were identified as 21 β -tigloyl-22 α -acetylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1–2)] [β -D-glucopyranosyl (1–4)]- β -D-glucopyranosiduronic acid (escin Ia, **1**), 21 β -angeloyl-22 α -acetylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1–2)] [β -D-glucopyranosyl (1–4)]- β -D-glucopyranosiduronic acid (escin Ib, **2**), 21 β -tigloyl-28-acetylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1–2)] [β -D-glucopyranosyl (1–4)]- β -D-glucopyranosiduronic acid (isoescin Ia, **3**), and 21 β -angeloyl-28-acetylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1–2)] [β -D-glucopyranosyl (1–4)]- β -D-glucopyranosiduronic acid (isoescin Ib, **4**) by comparison of their spectral data with the literature data.^{4–6} A single-crystal X-ray analysis undertaken on escin Ia* supported the structure elucidated based on NMR evidences and further suggested its absolute configuration (Figures 1 and 2).

Compound **5** was isolated as an amorphous powder, with a compositions of C₅₅H₈₆O₂₄ as determined by negative-ion

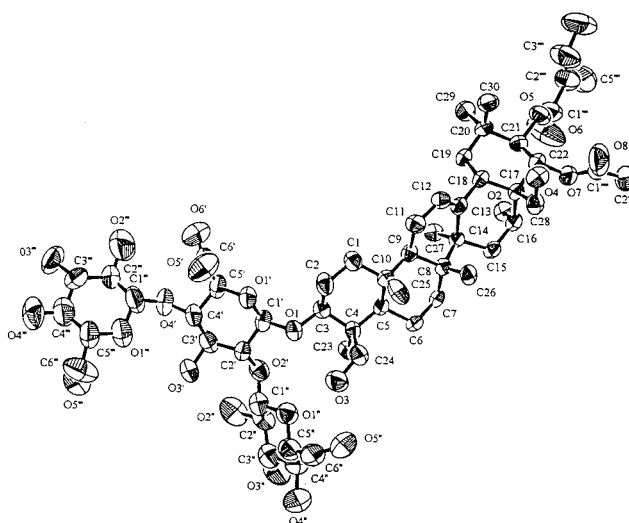


Figure 1. ORTEP drawing with atomic numbering of escin Ia.

HRSIMS, identical with **1–4**. The ¹H and ¹³C NMR signals of **5** showed a close resemblance to those of **1** and **3**,^{4,6} with presences of a protoaescigenin skeleton, a trisaccharide moiety, a tigloyl group, and an acetyl group. The significant differences in ¹³C and ¹H NMR spectra from those of **1** and **3** were chemical shifts of C-21 (δ 76.2) and C-22 (δ 77.9) with corresponding protons at δ 4.95 and 5.92. Otherwise, minor changes were also observed for C-17 (δ 45.9), C-18 (δ 41.4), and H-18 (δ 2.75). Compared with **1–4**,^{4,6} the stereochemistry of C-21 and C-22 remained unchanged as deduced from NOESY spectrum: Me-29 showed a strong NOE correlation with H-21 while Me-30 correlated with H-22 strongly. These indicated the α -configuration of H-21 and the β -configuration of H-22. The coupling constant of H-21 and H-22 ($J = 9.5$ Hz) also supported the proposed structure. The attachments of the tigloyl group at C-22 and the acetyl group at C-28 were derived from a HMBC experiment, which correlated the carbonyl carbons of the tigloyl and the acetyl groups to H-22 and H-28, respectively. ¹H and ¹³C NMR signals of the trisaccharide moiety were coincident with those of **1–4**, and acid hydrolysis of **1** also

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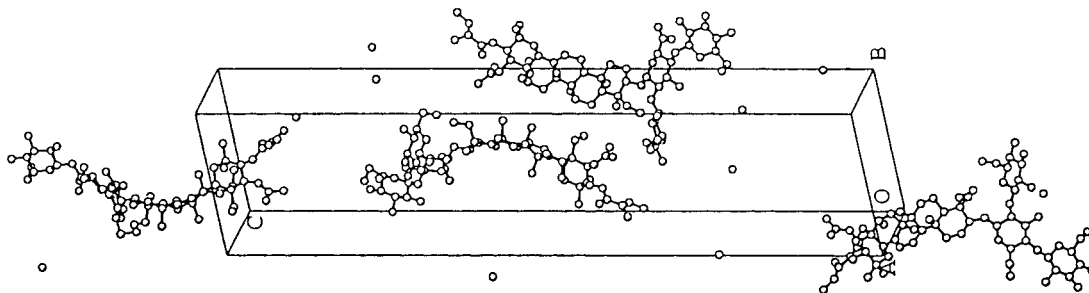


Figure 2. Stereoview of the unit-cell packing of escin Ia.

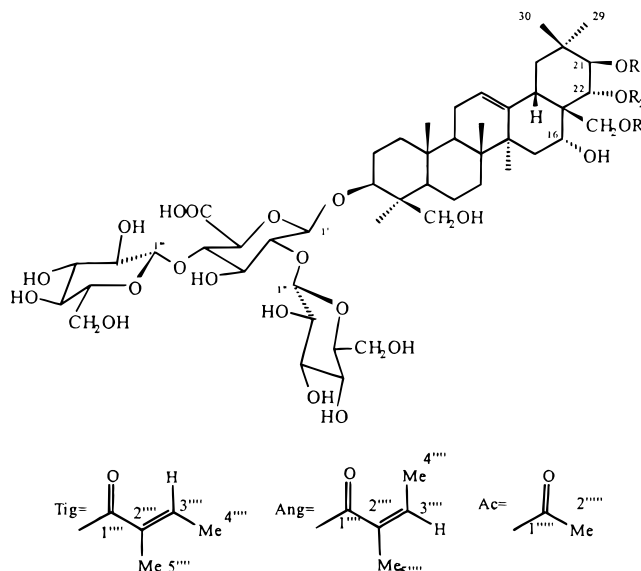
yielded glucose and glucuronic acid. Furthermore, their sequences and linkage sites were confirmed by HMBC correlations between the following pairs: C-3 (δ 90.9) and H-1' (δ 4.87); C-2' (δ 79.5) and H-1'' (δ 5.58); and C-4' (δ 81.7) and H-1''' (δ 5.18). Hence, **5** was established as 22 α -tigloyl-28-acetylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1-2)] [β -D-glucopyranosyl (1-4)]- β -D-glucopyranosiduronic acid and named escin IVc.

Compound **6** was isolated as an amorphous powder. It was determined to be an isomer of **5** by HRMALDIMS with the molecular formula of C₅₅H₈₆O₂₄. The only differences in their ¹H and ¹³C NMR spectra were signals due to an angeloyl group [¹³C NMR: δ 168.1 (C-1'''), 129.1 (C-2'''), 136.6 (C-3'''), 15.7 (C-4'''), and 20.6 (C-5''')]. ¹H NMR: δ 5.83 (H-3'''), 1.43 (Me-4'''), and 1.92 (Me-5''')] replaced those due to a tigloyl group. Similar HMBC and NOE correlations were also present for **6**. Therefore, **6** was identified as 22 α -angeloyl-28-acetylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1-2)] [β -D-glucopyranosyl (1-4)]- β -D-glucopyranosiduronic acid, and named escin IVd. It is a geometrical isomer of **5** (Figure 3).

Compound **7** was isolated as an amorphous powder. It has a composition of C₅₃H₈₄O₂₃ as shown by HRMALDIMS. Compared with **1-4** and **5**, ¹H and ¹³C NMR signals belonging to an acetyl group disappeared. Long-range correlation between the carbonyl carbon of the tigloyl group and H-28 (δ 4.15) characterized the presence of a tigloyl group at C-28. Otherwise, ¹H and ¹³C NMR signals due to the sugar moiety remained unchanged. On the basis of the evidence above, **7** was established as 28-tigloylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1-2)] [β -D-glucopyranosyl (1-4)]- β -D-glucopyranosiduronic acid and named escin IVe.

Compound **8** was isolated as an amorphous powder. Compounds **7** and **8** were determined to be isomeric, with a composition of C₅₃H₈₄O₂₃ as shown by negative-ion HRSIMS. Compared with **7**, ¹H and ¹³C NMR signals assignable to an angeloyl group [¹³C NMR: δ 167.6 (C-1'''), 128.0 (C-2'''), 138.1 (C-3'''), 15.8 (C-4'''), and 20.8 (C-5''')]. ¹H NMR: δ 5.84 (H-3'''), 1.94 (Me-4'''), and 1.85 (Me-5''')] replaced those due to a tigloyl group. The position of the angeloyl group and the sequence of the trisaccharide were consistent with **7** as determined by HMBC spectrum. Acid hydrolysis of **8** yielded glucose and glucuronic acid. On the basis of the above evidences, **8** was established as 28-angeloylprotoaescigenin-3 β -O-[β -D-glucopyranosyl (1-2)] [β -D-glucopyranosyl (1-4)]- β -D-glucopyranosiduronic acid and named escin IVf. Compounds **7** and **8** are also geometrical isomers.

Escins inhibited 86.1 \pm 0.2% of the HIV-1 protease activity at 100 μ M. The inhibition percentages of mixtures of its main components, **1+2** (2:1) and **3+4** (2:1), were 89.9 \pm 1.1% and 50.8 \pm 0.4% at 100 μ M. Of the isolated compounds (**1-8**) that were tested, compounds **1** and **2** showed inhibitory activity against HIV-1 protease with IC₅₀ values of 35 and 50 μ M, respectively. The inhibition



	R ₁	R ₂	R ₃
1	Tig	Ac	H
2	Ang	Ac	H
3	Tig	H	Ac
4	Ang	H	Ac
5	H	Tig	Ac
6	H	Ang	Ac
7	H	H	Tig
8	H	H	Ang

Figure 3. Structures of compounds **1-8**.

percentages of **3-8** at 100 μ M were 39.0 \pm 4.4%, 14.7 \pm 0.7%, 35.2 \pm 2.3%, 34.0 \pm 9.4%, 15.9 \pm 0.5%, and 12.5 \pm 1.7%, respectively. This is the first report of their anti-HIV-1-protease activity.

Experimental Section

General Experimental Procedures. Melting points are uncorrected. IR studies were made with KBr disks. NMR studies employed a Varian INOVA-500 Spectrometer operating at 500 MHz for ¹H and 125 MHz for ¹³C including COSY, HMQC, HMBC, and NOESY, and chemical shifts are given in δ relative to TMS as internal standard. MALDIMS (matrix-assisted laser desorption ionization MS) was taken on BIFLEX III (Bruker) and SIMS was taken on APEX II (Bruker). For preparative HPLC (pump, P2000; detector, UV 3000; software, PC 1000; Thermo Separation Products, U.S.A.) separation, an ODS column [Phenomenex LUNA 10 μ C₁₈ (2) (250 \times 21.2 i.d.), U.S.A.] column was used. A YMC-Pack ODS-AP (4.6 \times 150 mm, YMC, Kyoto, Japan) was used for anti-HIV-1 protease assay. The integrator was a C-R1A Chromatopac (Shimadzu, Japan) product.

5'''), 1.94 (3H, d, $J = 7.0$ Hz, Me-4'''), 2.84 (1H, d, $J = 14.0$ Hz, H-18), 2.97 (1H, dd, $J = 12.0, 13.0$ Hz, H-19b), 3.20 (1H, d, $J = 10.0$ Hz, H-24a), 3.26 (1H, d, $J = 11.0$ Hz, H-3 α), 3.58 (1H, d, $J = 8.5$ Hz, H-6''a), 4.12 (1H, d, $J = 8.0$ Hz, H-28a), 4.15 (1H, d, $J = 10.5$ Hz, H-24b), 4.22 (1H, d, $J = 11.0$ Hz, H-6''b), 4.35 (1H, d, $J = 9.0$ Hz, H-22 β), 4.38 (1H, d, $J = 8.5$ Hz, H-28b), 4.71 (1H, br s, H-16 β), 4.74 (1H, d, $J = 9.0$ Hz, H-21 α), 4.75 (1H, d, $J = 7.0$ Hz, H-1'), 5.10 (1H, d, $J = 7.5$ Hz, H-1'''), 5.38 (1H, br s, H-12), 5.56 (1H, d, $J = 7.5$ Hz, H-1''), 5.84 (1H, dq-like, H-3'''). ^{13}C NMR data, see Table 1; negative-ion HRSIMS m/z $[\text{M} - \text{H}^+]^-$ 1087.5319 (calcd for $\text{C}_{83}\text{H}_{83}\text{O}_{23}$, 1087.5330).

HIV-Protease Assay. A 25- μL amount of HIV-PR assay buffer (50 mM NaOAc, pH 4.9) containing 2.5 μg of substrate [His-Lys-Ala-Arg-Val-Leu-($p\text{NO}_2$ -Phe)-Glu-Ala-Nle-Ser-NH $_2$] was mixed with 2.5 μL of a compound (**1–8**) solution (using DMSO as solvent), and then 6.25 μL of recHIV-PR (0.25 μg of protein from Bachem) was added into this mixture. The reaction mixture was incubated for 15 min at 37 $^\circ\text{C}$ and then terminated by addition of 2.5 μL of 10% trifluoroacetic acid (TFA). The hydrolysate ($p\text{NO}_2$ -Phe-Glu-Ala-Nle-Ser-NH $_2$) and remaining substrate were quantitatively analyzed by reversed-phase HPLC with a gradient of acetonitrile (20–40% in 26 min) in 0.1% TFA at a flow rate of 1.0 mL/min. The elution profile was monitored at 280 nm. The substrate and the hydrolysate were eluted at 8.1 and 3.5 min, respectively. The peak areas were measured with an integrator. The inhibitory activity of a compound of HIV-PR was calculated as follows: %inhibition = $(A_{\text{control}} - A_{\text{sample}}) \times 100/A_{\text{control}}$ (where A is a

relative peak area of the hydrolysate). Acetylpepstatin was used as a positive control, $\text{IC}_{50} = 0.30 \mu\text{M}$.

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Supporting Information Available: Tables of X-ray data for escin Ia, including hydrogen bonds, atomic parameters, bond lengths and angles, and anisotropic parameters. This material is available free of charge via the Internet at <http://pubs.acs.org>. Crystallographic data for the structure reported in this paper have been deposited with the Cambridge Crystallographic Data Centre. Copies of the data can be obtained, free of charge, on application to the Director, CCDC, 12 Union Road, Cambridge CB2 1EZ, UK (Fax: +44 (0)1223-336033 or e-mail: deposit@ccdc.cam.ac.uk).

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