



Amyloid β aggregation inhibitory activity of triterpene saponins from the cactus *Stenocereus pruinosus*

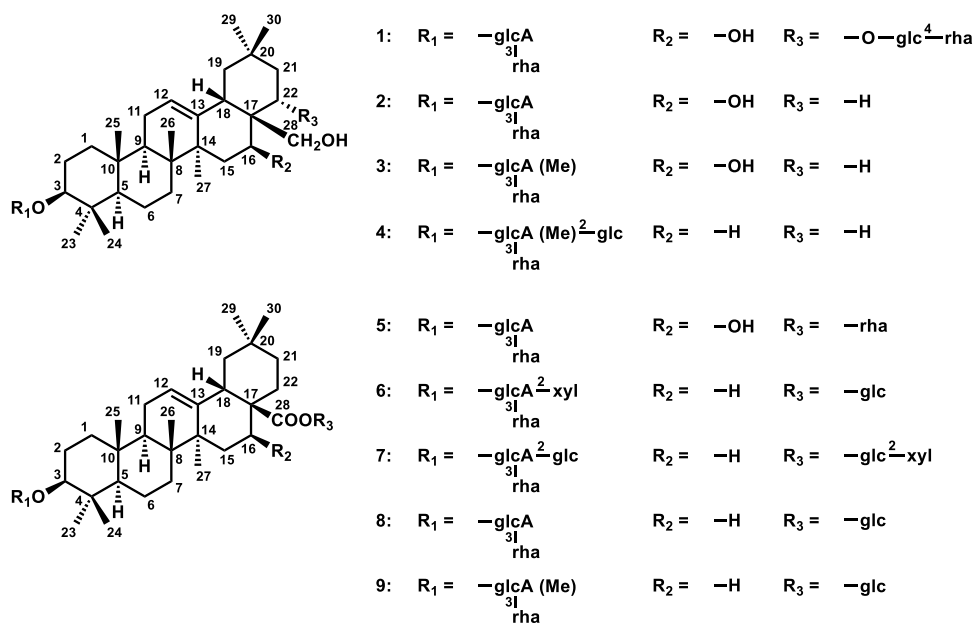
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Abstract

Six new triterpene saponins (1–5,7) and 3 known saponins (6,8,9) were isolated from MeOH extracts of the cactus *Stenocereus pruinosus*. The structures of the isolated saponins were elucidated using MS, IR, and comprehensive NMR measurements. To develop drugs for treating Alzheimer's disease (AD) on the basis of the amyloid cascade hypothesis, the isolated saponins were evaluated for inhibition of BACE1 activity and amyloid beta (A β) aggregation using thioflavin-T assay, and triterpenes as an aglycone moiety and an alkaline hydrolysate of the saponins were also evaluated. One saponin, stenoside A (7), exhibited inhibitory activity related to A β aggregation and its degree of A β aggregation was 40.6% at 100 μ M.

Graphic abstract



Keywords Cactus · *Stenocereus pruinosus* · Saponins · Alzheimer's disease · Amyloid beta · BACE1

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Introduction

Cacti range mainly throughout South America to the southern areas of North America, regions that have two seasons clearly distinguishable as rainy and dry. Cactus plants are

divided into primarily three forms, known as arborescent cacti, columnar cacti, and globular cacti. *Stenocereus pruinosus* (Otto) Buxb. belongs to columnar cacti widely distributed in semi-arid areas of the South East of Mexico, intensely managed in Central Mexico to gather its fruits, and sometimes cultivated as home garden [1]. The height of *S. pruinosus* is up to 8 m, blanches are green with 5–8 ribs, and flowers are infundibuliform 7–10 cm in length growing in the blanch apexes with green-brownish external tepals and white internal tepals which are produced 2 or 3 years after being planted [1, 2]. The constituents of cacti have been investigated by Djerassi and co-workers, who reported a lot of triterpenoid saponin in an acid-hydrolyzed saponin-rich extract from many cacti and one of their works revealed *S. pruinosus* contained oleanolic acid [3]. Considering those reports, we had been further investigating triterpene saponin from many cacti and discovered that *S. pruinosus* contained erythrodiol, longispinogenin and 3 β -hydroxy-11 α ,12 α -epoxyolean-28,13 β -olide in addition to oleanolic acid from hydrolysate of MeOH extract of *S. pruinosus* [4]. Now, we have been investigating triterpene saponin from cacti for several decades [5–12] and reported the identification of numerous saponin exhibiting bioactivities such as anti-type I allergy activity [7], inhibition or promotion of melanin synthesis [5], inhibition of amyloid β (A β) aggregation, and protective effects on SH-SY5Y cells against A β -associated toxicity [6].

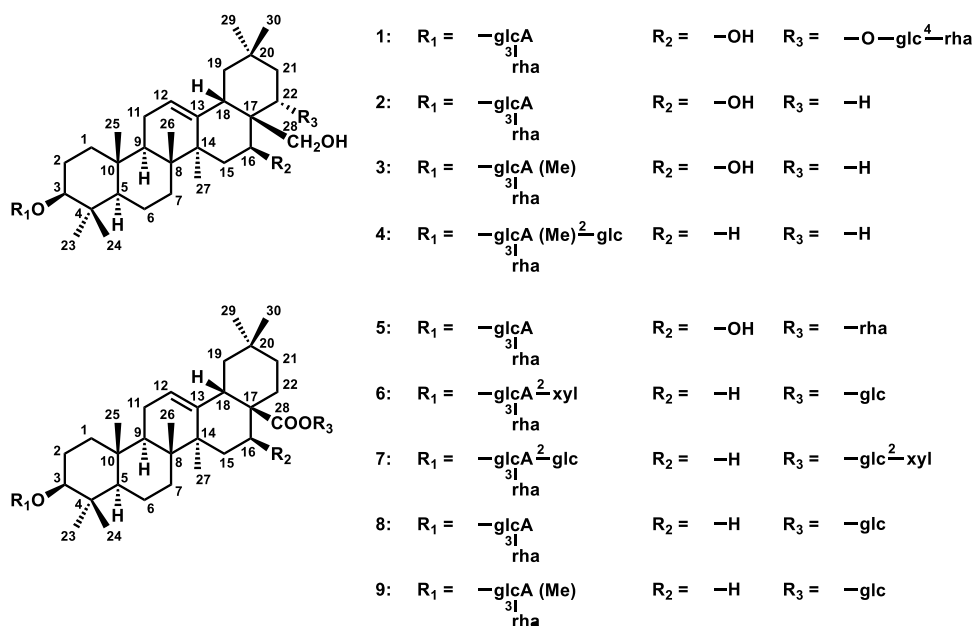
Saponin are well-known phytochemicals that are comprised of two parts, an aglycone moiety and a sugar moiety. The activities of saponin have been well studied and include anti-inflammatory, anti-tumor, anti-obesity, anti-angiogenesis, anti-allergic, anti-microbial, and anti-Alzheimer's disease (AD) [13, 14]. In particular, our group found that

saponin from cacti have unique structures [5–12] that have not been adequately investigated.

AD is a globally significant disease that involves cognitive and locomotor disorders resulting from progressive neurodegeneration. It is frequently said that AD has a complex pathogenesis that involves several key steps: A β production, A β aggregation, and neural cell death. A β production is caused by the cleavage of amyloid precursor protein (APP) by the action of the enzymes β -secretase and γ -secretase [15, 16]. As the concentration of A β increases, it begins to aggregate and form A β -oligomers, A β -fibrils, and in most cases, senile plaques. Once A β aggregation occurs, neural cell death gradually begins [17, 18]. This step also involves complex pathways that consist of synaptic and neuritic injury, altered ionic homeostasis, oxidative damage, altered kinase/phosphatase activity, neurofibrillary tangles, and microglial and astrocytic activation [19]. The prevention of neural cell death is clearly important, but A β production and aggregation are also important for the A β -associated steps that represent the upstream pathways of AD pathogenesis. As such, many groups have explored numerous candidate phytochemicals from natural resources for inhibition of A β production and aggregation [19–22]. A recent report identified saponin as a class of candidate phytochemicals [23, 24].

Here, we report the isolation and structure elucidation of various saponin (1–9), including 6 new compounds (1–5,7) and 3 known saponin (6,8–9), isolated from the cactus *Stenocereus pruinosus* (Otto) Buxb. (Fig. 1). We also evaluated the activity of these compounds in terms of inhibition of A β aggregation and BACE1 activity. Our data revealed two particularly remarkable findings. First, compounds 1 and 6–7 have a rare linkage of a sugar unit. Compound 1

Fig. 1 Structures of compounds (1–9) Isolated from *Stenocereus pruinosus*



has a glucopyranosyl unit that binds to the C-22 region of its aglycone, whereas **6** has a xylopyranosyl unit that binds to the C-3' of glucuronic acid, and **7** also has a xylopyranosyl unit that binds to the C-2'' of glucose binding to C-28 of its aglycone. Second, compounds **2–4** have rare triterpenes for cactus plants, longispinogenin and erythrodiol, as aglycones. These are the first reports of isolation of such saponins from columnar cactus plants.

Results and discussion

Dried *Stenocereus pruinosus* was extracted repeatedly with chloroform and then with methanol. The methanol extract was separated by silica gel and octadecyl silyl silica gel (ODS) column chromatography, yielding chichipenoside D (**1**), longispinoside A (**2**), longispinoside A methyl ester (**3**), erythronoside A (**4**), cochalinoside C (**5**), oleanolic acid 3-*O*- β -D-xylopyranosyl-(1 \rightarrow 2)- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranosyl-28-*O*- β -D-glucopyranoside (**6**), Stenoside A (**7**), oleanolic acid 3-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranosyl-28-*O*- β -D-glucopyranoside (**8**) and oleanolic acid 3-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-6'-*O*-methyl glucuronopyranosyl-28-*O*- β -D-glucopyranoside (**9**) (Fig. 1). The structures of compounds **1–9** were determined by mass and NMR spectroscopy using ^1H - and ^{13}C -NMR, DEPT, HMQC, HMBC, ^1H - ^1H COSY or DQF-COSY, HSQC-TOCSY, phase-sensitive TOCSY and phase-sensitive NOESY experiments.

Compound **1** (16.7 mg, 0.003%) was obtained as a colorless amorphous powder, and its optical rotation value was elucidated as $[\alpha]_D^{21}$ -28.0 (*c* 0.25, MeOH). The molecular formula of **1** was elucidated to be $\text{C}_{54}\text{H}_{88}\text{O}_{23}$ using negative HRFABMS (*m/z* 1103.5640, $[\text{M}-\text{H}]^-$). The IR spectrum of **1** showed absorptions at 3385 cm^{-1} (hydroxy) and 1714 cm^{-1} (carbonyl). The signal patterns of the ^1H - and ^{13}C -NMR spectral data indicated that the aglycone moiety

was an oleanane-type triterpene. The ^1H - and ^{13}C -NMR spectral data indicated the presence of seven methyl groups in the aglycone, characterized by signals at $[\delta_{\text{H}} 0.76, 0.87, 0.90, 0.96$ and 1.16 (each 3H, s), and $\delta_{\text{H}} 0.88$ (6H, s)], and $15.3, 16.4, 16.6, 24.4, 26.8, 27.6,$ and 32.8 (each CH_3). The presence of signals at $\delta_{\text{C}} 64.5$ (CH) and $\delta_{\text{H}} 4.42$ (1H, br s), $\delta_{\text{C}} 77.9$ (CH) and $\delta_{\text{H}} 4.25$ (1H, dd, $J = 12.6, 4.5$ Hz), $\delta_{\text{C}} 56.7$ (CH_2) and $[\delta_{\text{H}} 3.33$ (1H, o), and $\delta_{\text{H}} 3.77$ (1H, m)] indicated that the aglycone had three hydroxyl groups, except in the C-3 region [$\delta_{\text{C}} 88.1$ (CH) and $\delta_{\text{H}} 3.02$ (1H, dd, $J = 11.3, 3.2$ Hz)], and the stereochemistry of 3-OH was determined as a β -configuration by its $^3J_{\text{H}_2/\text{H}_3}$ coupling constants of 11.3 Hz. Further analysis of the aglycone of **1** primarily using HMQC, HMBC, DQF-COSY, phase-sensitive NOESY spectral correlations, and complementarily using phase-sensitive TOCSY spectral correlations, revealed four hydroxyl groups assigned at C-3 ($\delta_{\text{C}} 88.1$), C-16 ($\delta_{\text{C}} 64.5$), C-22 ($\delta_{\text{C}} 77.9$), and C-28 ($\delta_{\text{C}} 56.7$). The NOESY correlations between $\delta_{\text{H}} 1.16$ (H_3 -27) and $\delta_{\text{H}} 4.42$ (H-16), $\delta_{\text{H}} 3.33, 3.77$ (H-28) and $\delta_{\text{H}} 0.90$ (H-26), and $\delta_{\text{H}} 4.25$ (H-22) and $\delta_{\text{H}} 2.50$ (H-18) supported the stereochemistries of 16-OH and 22-OH as β -configuration and α -configuration, respectively (Fig. 2). The signals at $\delta_{\text{H}} 1.56$ (1H, br t, $J = 12.6$ Hz, H-21 α), $\delta_{\text{H}} 1.68$ (1H, br d, $J = 12.6$ Hz, H-21 β) and $\delta_{\text{H}} 4.25$ (1H, dd, $J = 12.6, 4.5$ Hz, H-22) also supported the stereochemistry of 22-OH as α -configuration by its large $^3J_{\text{H}_{21\alpha}/\text{H}_{22}}$ coupling constants of 11.3 Hz and small $^3J_{\text{H}_{21\beta}/\text{H}_{22}}$ coupling constants of 4.5 Hz. These data confirmed that the aglycone of **1** was chichipegenin. The ^1H - and ^{13}C -NMR spectra exhibited four anomeric proton signals [$\delta_{\text{H}} 4.17$ (1H, d, $J = 7.7$ Hz), $\delta_{\text{H}} 4.33$ (1H, d, $J = 7.8$ Hz), $\delta_{\text{H}} 4.69$ (1H, br s), $\delta_{\text{H}} 5.04$ (1H, br s)] and carbon signals [$\delta_{\text{C}} 100.2$ (CH), $\delta_{\text{C}} 100.5$ (CH), $\delta_{\text{C}} 103.3$ (CH), $\delta_{\text{C}} 105.1$ (CH)]. Detailed analysis of the sugar moiety of **1** using HMQC, HMBC, DQF-COSY, HSQC-TOCSY, phase-sensitive TOCSY, and phase-sensitive NOESY experiments revealed the presence of one β -glucuronic acid, one β -glucose, and two α -rhamnose units. The linkage of each sugar was determined by HMBC correlations.

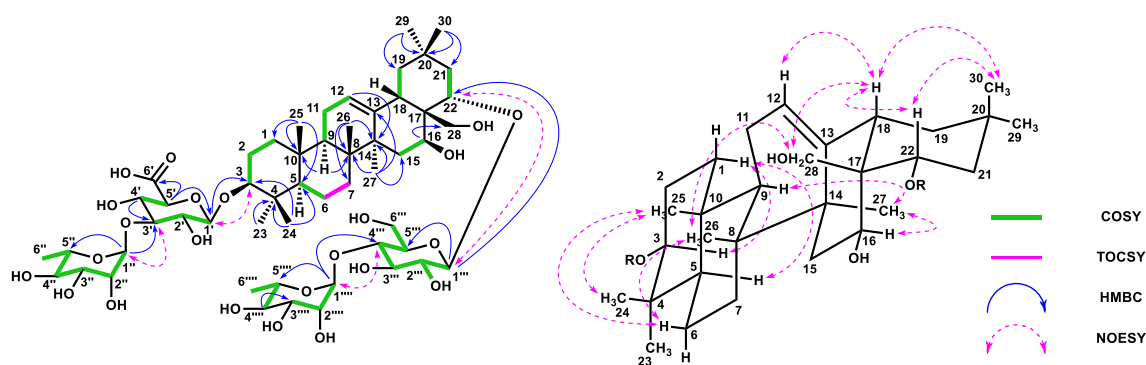


Fig. 2 HMBC, COSY, and Key NOESY correlations of **1**

The correlation between H-1' of the glucuronic acid (δ_{H} 4.17) and C-3 of the aglycone (δ_{C} 88.1) showed that the glucuronic acid-1 unit binds at C-3 of the aglycone moiety. Similarly, the correlation between H-1'' of rhamnose (δ_{H} 5.04) and C-3' of glucuronic acid (δ_{C} 80.3) showed that the rhamnose-1 unit binds at C-3' of the glucuronic acid unit. In contrast, the correlation between H-1''' of glucose (δ_{H} 4.33) and C-22 of the aglycone (δ_{C} 77.9) showed that the glucose-1 unit binds at C-22 of the aglycone moiety. Considering the chemical shift of C-22 (δ_{C} 67.4) of chichipenoside C [5], this deshielding of C-22 (δ_{C} 77.9) of the aglycone also supported the binding of the glucose-1 unit at C-22 of the aglycone. In addition, the correlation between H-1'''' of the second rhamnose (δ_{H} 4.69) and C-4''' of glucose (δ_{C} 76.6) indicated that the rhamnose-1 unit binds at C-4''' of the glucose unit. These correlations are shown in Fig. 2. The absolute configuration of each sugar unit was determined by measuring the ODS HPLC retention times of the derivatives from D-glucuronic acid, D-glucose, and L-rhamnose [25]. Whereas the anomeric proton on the glucuronopyranosyl and glucopyranosyl units were determined to have a β -configuration based on the $^3J_{\text{H}_1\text{H}_2}$ values (7.7 and 7.8 Hz, respectively), the anomeric protons on the two rhamnopyranosyl units were determined to have a α -configuration based on $^1J_{\text{C}_1\text{H}_1}$ value (171.7 and 169.4 Hz) [26] and three-bond strong HMBC correlations from H-1'' to C-3'' and C-5'', and H-1'''' to C-3'''' and C-5'''. Its three-bond strong HMBC correlations were attributed to the dihedral angles between H-1 and C-3, H-1 and C-5 of rhamnose about 180° [27]. Thus, compound **1** was determined chichipegenin 3-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranosyl-22-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 4)- β -D-glucopyranoside and named as chichipenoside D (Fig. 2).

Compound **2** (16.0 mg, 0.003%) was obtained as a colorless amorphous powder, and its optical rotation value was elucidated as $[\alpha]_{\text{D}}^{18} -52.0$ (*c* 0.2 MeOH). The molecular formula of **2** was elucidated to be $\text{C}_{42}\text{H}_{68}\text{O}_{13}$ using positive HRFABMS (m/z 803.4546, $[\text{M} + \text{Na}]^+$). The IR spectrum

of **2** showed absorptions at 3330 cm^{-1} (hydroxy) and 1740 cm^{-1} (carbonyl). The ^1H - and ^{13}C -NMR spectral data indicated the presence of seven methyl groups in its aglycone, characterized by signals at [δ_{H} 0.75, 0.87, 0.90, 0.96 and 1.14 (each 3H, s), and 0.84 (6H, s)] and 15.3, 16.3 16.5, 23.7, 26.6, 27.5, and 33.1 (each CH_3). The signal patterns of the ^1H - and ^{13}C -NMR spectral data indicated that the aglycone moiety was an oleanane-type triterpene. The presence of three sets of signals at δ_{C} 64.5 (CH) and δ_{H} 4.06 (1H, dd, $J = 12.4, 4.4$ Hz), δ_{C} 65.2 (CH_2) and δ_{H} 3.12 (1H, o), 3.51 (1H, d, $J = 10.2$ Hz) and δ_{C} 88.3 (CH) and δ_{H} 3.05 (1H, dd, $J = 11.3, 4.0$ Hz) indicated that the aglycone of **2** had three hydroxyl groups. Further analysis of the aglycone of **2** using HMQC, HMBC, DQF-COSY, and phase-sensitive NOESY spectral correlations assigned the three hydroxyl groups at C-3, C-16, and C-28. The NOESY correlation between δ_{H} 1.14 (H_3 -27) and δ_{H} 4.06 (H-16) and the $^3J_{\text{H}_{15}/\text{H}_{16}}$ coupling constant of 12.4 Hz revealed the stereochemistry of the 16-OH was β -configuration and confirmed that the aglycone of **2** was longispinogenin (Fig. 3). The ^1H - and ^{13}C -NMR spectra about sugar units showed partially similar signal patterns to those of **1**. Additionally considering ODS HPLC data [25] and comprehensive 2D-NMR analysis including strong three-bond HMBC correlation [27], it was revealed that the presence of two sugars, β -D-glucuronic acid and α -L-rhamnose. It could not be performed to measure the $^1J_{\text{C}_1\text{H}_1}$ value about the anomeric protons on the rhamnopyranosyl units contained in new saponins (**2–5,7**) from *S. pruinosis*, the rhamnopyranosyl units presumably had α -configuration because of our previous reports [5–12], their multiplicity (“br s” or “d” with small coupling constant) and their strong three-bond HMBC correlations [27]. Further analysis confirmed that compound **2** was longispinogenin 3-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranoside and named as longispinoside A (Fig. 3).

Compound **3** (24.2 mg, 0.004%) was obtained as a colorless amorphous powder, and its optical rotation value was elucidated as $[\alpha]_{\text{D}}^{19} -117.3$ (*c* 0.1 MeOH). The molecular

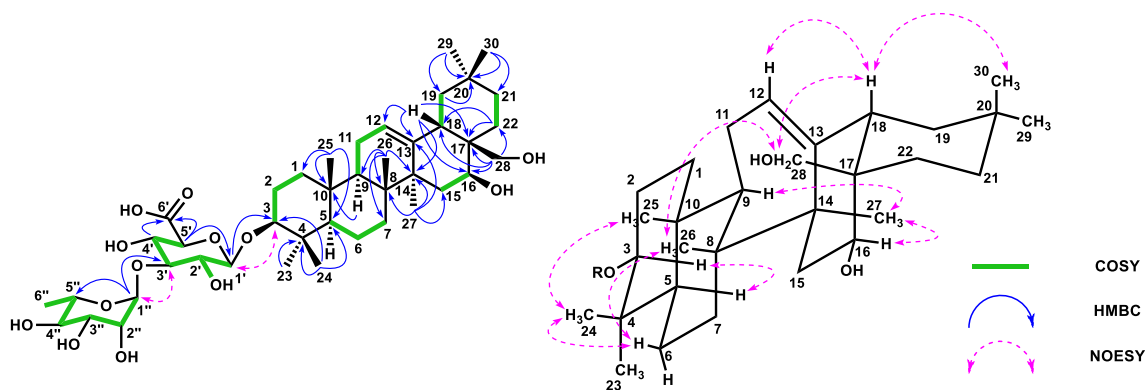


Fig. 3 HMBC, COSY, and Key NOESY correlations of **2**

formula of **3** was elucidated to be $C_{43}H_{70}O_{13}$ using positive HRFABMS (m/z 795.4900, $[M+H]^+$). The IR spectrum of **3** showed absorptions at 3363 cm^{-1} (hydroxy) and 1746 cm^{-1} (carbonyl). The ^1H - and ^{13}C -NMR spectral data were almost the same with **2** except for methyl signal [δ_{C} 51.9 (CH_3) and δ_{H} 3.65 (3H, s)] and C-6' of glucuronic acid [δ_{C} 169.5 (C)]. This differences and further analysis confirmed that compound **3** was longispinogenin 3-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranoside methyl ester and named as longispinoside A methyl ester (Fig. S1).

Compound **4** (9.3 mg, 0.001%) was obtained as a colorless amorphous powder, and its optical rotation value was elucidated as $[\alpha]_{\text{D}}^{19}$ -4.1 (c 0.1 MeOH). The molecular formula of **4** was elucidated to be $C_{49}H_{80}O_{17}$ using positive HRFABMS (m/z 941.5481, $[M+H]^+$). The IR spectrum of **4** showed absorptions at 3370 cm^{-1} (hydroxy) and 1743 cm^{-1} (carbonyl). The ^1H - and ^{13}C -NMR spectral data indicated the presence of seven methyl groups in its aglycone, characterized by signals at [δ_{H} 0.74, 0.82, 0.84, 0.95 and 1.10 (each 3H, s), and 0.87 (6H, s)], and 15.3, 16.1, 16.4, 23.5, 25.6, 27.5, and 33.1 (each CH_3). The signal patterns of the ^1H - and ^{13}C -NMR spectral data indicated that the aglycone moiety was an oleanane-type triterpene. The presence of two sets of signals at [δ_{C} 67.4 (CH_2) and δ_{H} 2.93 (1H, o) and 3.29 (1H, dd, $J=9.6, 4.8\text{ Hz}$)] and [δ_{C} 88.6 (CH) and δ_{H} 3.04 (1H, dd, $J=11.3, 4.0\text{ Hz}$)] indicated that the aglycone of **4** had two hydroxyl groups in the C-3 region. Further analysis of the aglycone of **4** using HMQC, HMBC, DQF-COSY, phase-sensitive TOCSY, and phase-sensitive NOESY spectral correlations assigned the two hydroxy groups at C-3 and C-28, which indicated that the aglycone of **4** was erythrodiol (Fig. 4). The ^1H - and ^{13}C -NMR spectra showed three anomeric proton signals [δ_{H} 4.43 (1H, d, $J=7.9\text{ Hz}$), δ_{H} 4.52 (1H, d, $J=8.3\text{ Hz}$), δ_{H} 4.94 (1H, br s)] and carbon signals [δ_{C} 100.5 (CH), δ_{C} 101.6 (CH), δ_{C} 103.1 (CH)].

Detailed analysis of the sugar moiety of **4** using HMQC, HMBC, DQF-COSY, HSQC-TOCSY, phase-sensitive TOCSY, and phase-sensitive NOESY experiments revealed the presence of one β -glucuronic acid, one β -glucose, and one α -rhamnose unit, and the signals at δ_{H} 3.65 (3H, s) and δ_{C} 169.5 indicated the presence of glucuronic acid methyl ester instead of glucuronic acid. The linkage of each sugar was determined by HMBC correlations and the correlations indicated that the glucuronic acid-1 unit binds at C-3 of the aglycone moiety, the glucose unit binds at C-2' of the glucuronic acid methyl ester, and the rhamnose-1 unit binds at C-3' of the glucuronic acid methyl ester (Fig. 4). The sugar units were determined as β -D-glucuronic acid, α -L-rhamnose, and β -D-glucose by the same method as compound **2** [25, 27]. Thus, compound **4** was determined erythrodiol 3-*O*- β -D-glucopyranosyl-(1 \rightarrow 2)-[α -L-rhamnopyranosyl-(1 \rightarrow 3)]- β -D-glucuronopyranoside methyl ester and named as erythronoside A methyl ester (Fig. 4).

Compound **5** (27.5 mg, 0.004%) was obtained as a colorless amorphous powder, and its optical rotation value was elucidated as $[\alpha]_{\text{D}}^{22}$ -28.8 (c 0.25 MeOH). The molecular formula of **5** was elucidated to be $C_{48}H_{76}O_{18}$ using negative HRFABMS (m/z 939.4952, $[M-H]^-$). The IR spectrum of **5** showed absorptions at 3370 cm^{-1} (hydroxy) and 1710 cm^{-1} (carbonyl). The ^1H - and ^{13}C -NMR spectral data indicated the presence of seven methyl groups in its aglycone, characterized by signals at δ_{H} 0.69, 0.75, 0.84, 0.90, 0.92, 0.95, and 1.13 (each 3H, s) and δ_{C} 15.1, 16.4, 16.8, 23.7, 26.7, 27.5, and 32.7 (each CH_3). The signal patterns of the ^1H - and ^{13}C -NMR spectral data indicated that the aglycone moiety was an oleanane-type triterpene. The presence of a signal at δ_{C} 62.7 (CH) and δ_{H} 4.06 (1H, t-like, $J=11.8\text{ Hz}$) indicated that the aglycone of **5** had one hydroxyl group excepting the C-3 region [δ_{C} 88.1 (CH) and δ_{H} 3.03 (1H, dd, $J=9.9, 3.8\text{ Hz}$)]. Further analysis of the aglycone of **5** using HMQC, HMBC,

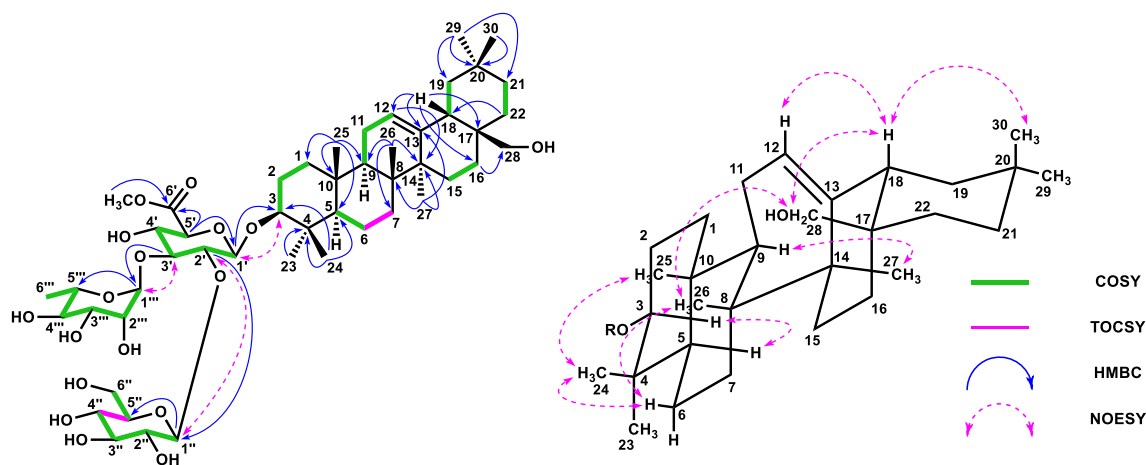


Fig. 4 HMBC, COSY, and Key NOESY correlations of **4**

DQF-COSY, phase-sensitive TOCSY, and phase-sensitive NOESY spectral correlations assigned the two hydroxy groups at C-3 and C-16. The stereochemistry of 16-OH was revealed as β -configuration by NOESY correlations and its $^3J_{H_{15}/H_{16}}$ coupling constant of 12.6 Hz, which indicated the aglycone of **5** was cochalic acid (Fig. 5). The ^1H - and ^{13}C -NMR spectra further indicated the presence of the same sugar moiety with **2** and another anomeric signals [δ_{H} 5.74 (1H, d, 1.1) and δ_{C} 93.3 (CH)]. Further analysis revealed the presence of another rhamnose unit binding at C-28 of the aglycone. The sugar units were determined as β -D-glucuronic acid and α -L-rhamnose by the same method as compound **2** [25, 27]. Thus, compound **5** was determined cochalic acid 3-O- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranosyl-28-O- α -L-rhamnopyranoside and named as cochalinoside C (Fig. 5).

Compound **7** (28.9 mg, 0.005%) was obtained as a colorless amorphous powder, and its optical rotation value was elucidated as $[\alpha]_{\text{D}}^{19}$ -8.6 (*c* 0.1 MeOH). The molecular formula of **7** was elucidated to be $\text{C}_{59}\text{H}_{94}\text{O}_{27}$ using positive HRFABMS (*m/z* 1273.5620, $[\text{M} + \text{K}]^+$). The IR spectrum of **7** showed absorptions at 3372 cm^{-1} (hydroxy) and 1732 cm^{-1} (carbonyl). The ^1H - and ^{13}C -NMR spectral data indicated that the aglycone was oleanolic acid, the presence of the same sugar moiety with **4** except for methyl ester, and the presence of two other sugar units. Further analysis revealed the presence of one glucose unit binding at C-28 of the aglycone and one xylose unit binding at C-2'''' of the glucose unit (Fig. S3). The sugar units were determined as β -D-glucuronic acid, α -L-rhamnose, β -D-glucose and β -D-xylose by the same method as compound **2** [25, 27]. Thus, compound **7** was determined oleanolic acid 3-O- β -D-glucopyranosyl-(1 \rightarrow 2)-[α -L-rhamnopyranosyl-(1 \rightarrow 3)]- β -D-glucuronopyranosyl-28-O- β -D-xylopyranosyl-(1 \rightarrow 2)- β -D-glucopyranoside and named as stenoside A (Fig. S3).

Compound **6** was analyzed in the same way as described above and elucidated as known one, oleanolic acid 3-O- β -D-xylopyranosyl-(1 \rightarrow 2)- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-

glucuronopyranosyl-28-O- β -D-glucopyranoside [28]. The 2D-correlations and completed assignment were shown in Fig. S2 and Table S1.

Compounds **8** and **9** were identified as oleanolic acid 3-O- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranosyl-28-O- β -D-glucopyranoside (**8**) and oleanolic acid 3-O- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-6'-O-methyl glucuronopyranosyl-28-O- β -D-glucopyranoside (**9**) by comparing their spectroscopic data with published data [29, 30].

Some of isolated saponins (**1–4, 7–8**), hydrolysate of **8** (**8a**), and oleanolic acid as the aglycone moiety of **6–9** were evaluated the inhibition of A β aggregation using a Th-T assay and inhibition of BACE1 activity (Table 1). A newly isolated saponin (**7**) exhibited A β aggregation inhibitory activity (40.6% at 100 μM), and compounds **1** and **4** exhibited moderate activity (66.4% and 70.0%, respectively, at 100 μM). The activity of **8a** was almost same level as **8** at 100 μM .

A preliminary SAR suggested that glycosylations at C-28 position of the oleanolic acid like **7** increased its inhibitory activity comparing with a glycosylation at C-28 position of the oleanolic acid like **8**. In addition to this, glycosylations at C-3 position of the oleanolic acid hardly had any effects on its activity in the case of **8a**. According to these results, glycosylations at C-28 might be important for A β aggregation inhibitory activity.

Chichipenoside C, chichipegenin 3-O- α -L-rhamnopyranosyl-(1 \rightarrow 2)- β -D-glucopyranosyl-(1 \rightarrow 2)- β -D-glucuronopyranoside, which has already isolated from *Polaskia chichipe* [5], exhibited A β aggregation inhibitory activity at the same level as chichipegenin (chichipenoside C; 43.8%, chichipegenin; 39.2%, respectively, at 25 μM) [6], while chichipenoside D exhibited weaker activity than that of chichipenoside C and chichipegenin. Its SAR suggested that glycosylations at C-22 position of the chichipegenin attenuated A β aggregation inhibitory activity. It would be expected that saponins mono-glycosylated at C-22 position

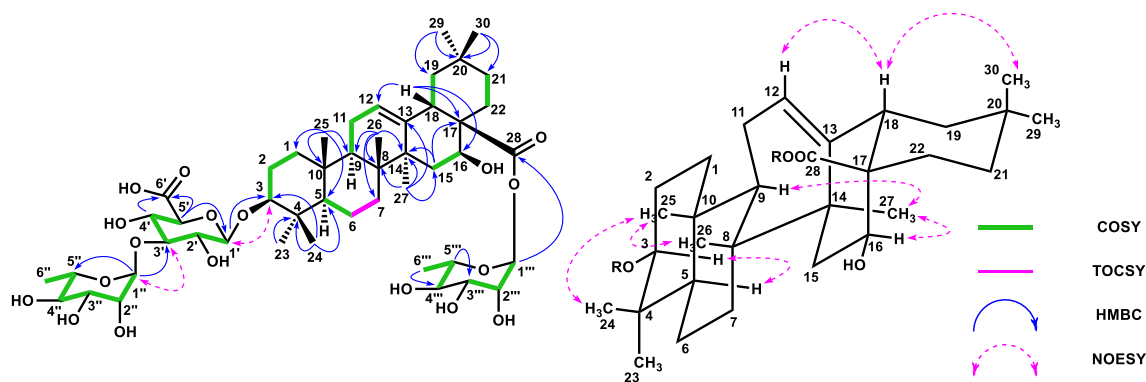


Fig. 5 HMBC, COSY, and Key NOESY correlations of **5**

Table 1 Assay results of the isolated and associated compounds against A β aggregation and BACE1 activity

Compound	Conc. (μ M)	A β aggregation (%)	BACE1 inhibition (%)
1	100	66.4 \pm 6.4	NA ^b
2	100	92.5 \pm 7.3	NA ^b
3	100	77.3 \pm 4.1	NA ^b
4	100	70.0 \pm 0.7	NA ^b
5	100	ND ^a	ND ^a
6	100	ND ^a	ND ^a
7	100	40.6 \pm 5.8	NA ^b
8	100	81.4 \pm 3.7	NA ^b
9	100	ND ^a	ND ^a
8a	100	95.4 \pm 15.4	NA ^b
Oleanolic acid	100	78.8 \pm 7.8	45.3 \pm 23.2
Cochalic acid	100	ND ^a	35.8 \pm 8.8
Chichipegenin	100	ND ^a	NA ^b
Myricetin	25	23.4 \pm 2.2	
Inhibitor IV	10		69.1 \pm 13.7

Data are shown as mean with standard errors ($n=3$)

^aND=no data

^bNA=not active

of the chichipegenin are isolated and tested for inhibitory activity related to A β aggregation.

In contrast, saponins (**1–4,7–8**) and hydrolysate of **8 (8a)** did not exhibit any BACE1 inhibitory activities at 100 μ M, while oleanolic acid exhibited moderate activity (45.3% at 100 μ M). Its IC₅₀ value was previously reported [31]. Considering oleanolic acid, cochalic acid and chichipegenin, the presence of a carboxyl group at C-28 region of the oleanane-type triterpenes might be effective for BACE1 inhibitory activity and the oleanane-type triterpenoid saponins might have less ability for inhibiting BACE1 activity.

In summary, six new saponins (**1–5,7**) were isolated from *Stenocereus pruinosus*, and their structures were elucidated along with known saponins (**6,8,9**). The structure of **1** has chichipegenin as the aglycone, in which C-3 and C-22 are glycosylated. Although saponins that have a sugar unit at C-22 to some extent have been identified [32–34], this is the first report in the case of chichipegenin. Compounds **2–3** and **4** have longispinogenin or erythrodiol as the aglycone moiety. Some saponins that have longispinogenin or erythrodiol as the aglycone have already been reported [35–42], but this is the first report of their isolation from cacti. Saponins **5** and **6–7** have cochalic acid or oleanolic acid as the aglycone. These types of saponin has been also isolated from another cactus, *Polaskia chichipe* [6], but the sugar unit differs from known ones. With respect to the sugar unit, **6** has a unique sugar chain for cactus plants. There are some reports related to that type of sugar chain [28, 43–47], as it is not a common

sugar unit, and this is the first report of its isolation from cacti. Paying attention to glucuronic acid, some compounds (**3, 4, 9**) were isolated as glucuronic acid methyl esters. As compound **3** and **9** appeared to be derivatives of **2** and **8**, compounds **3, 4** and **9** could be artifacts caused by methanol extraction, which thought to be similar case to our previous report [5]. Some of the isolated saponins (**1–4,7–8**) and hydrolysate of **8 (8a)** were tested for their ability to inhibit A β aggregation and BACE1 activity. Compound **7** was most effective at inhibiting A β aggregation, followed by compounds **1** and **4**. Oleanolic acid exhibited less of an effect on A β aggregation, as the structure consisting of oleanolic acid and sugar units is important for attenuating A β aggregation. Results observed with compound **7** also suggested that glycosylations at C-28 of oleanolic acid is important for attenuating A β aggregation. In the BACE1 assay, tested compounds exhibited no inhibitory effect on BACE1 activity except for oleanolic acid and cochalic acid. Although six new saponins were described, more isolations from cacti should be pursued because of their unique structures, and additional SARs should be conducted to characterize the relationship to inhibition of A β aggregation and BACE1 activity.

Experimental

General experimental procedures

Optical rotation values were recorded with a Horiba SEPA-300 polarimeter. IR spectra were measured with a Thermo FT-IR Nicolet iS5 spectrometer (ATR). ¹H- and ¹³C-NMR spectra were recorded using a JNM LA-500 spectrometer. HRFABMS spectra were obtained using a JEOL JMS-700 spectrometer. Column chromatography was carried out with silica gel 60 N (63–210 μ m) from Kanto Chemical, ODS silica gel YMC-GEL ODS-A from YMC Co. Ltd., and Diaion HP-20 from Mitsubishi Chemical Co. Ion-exchange chromatography was performed using Dowex 50 W-X8 resin (50–100 mesh, H-form) from Sigma-Aldrich. Thin-layer chromatography was carried out using TLC silica gel 60F₂₅₄ and RP-18 F_{254S} plates from Merck. Inhibition of A β aggregation was assayed using a Synergy HTX Multi-Mode Reader (BioTek, Winooski, Vermont, USA), A β ₄₀ (Peptide Institute, Osaka, Japan), and thioflavin-T (Fujifilm-Wako).

Plant material

Stenocereus pruinosus (Otto) Buxb. was purchased from Hokoan (Iga City, Mie, Japan). A voucher specimen is deposited at our laboratory.

Extraction and isolation

Aerial parts of *S. pruinosa* were dried, and the dry powder (639.1 g) was extracted three times with CHCl_3 and then extracted three times with MeOH. The MeOH extract (144.2 g) was applied to a Diaion HP-20 column, which was successively eluted with MeOH– H_2O mixture to give 4 fractions [H_2O -eluted fraction (disposal), 30% MeOH-eluted fraction (15.0 g), 70% MeOH-eluted fraction (60.3 g), and 100% MeOH-eluted fraction (106.5 g)], respectively.

The 70% MeOH-eluted fraction (60.3 g) was subjected to silica gel column chromatography (Si. C. C.) using a CHCl_3 –MeOH– H_2O mixture to give 10 fractions (A–J). Fr. H (10.3 g) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to give 3 fractions (Ha–Hc). Fr. Hb (7.5 g) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to give 5 fractions (Hb1–Hb5). Fr. Hb4 (5.0 g) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to give 3 fractions (Hb4-1–Hb4-3). Fr. Hb4-2 (4.1 g) was subjected to octadecyl-silylated silica gel column chromatography (ODS C. C.) using a MeOH– H_2O mixture to give 5 fractions (Hb4-2a–Hb4-2e). Fr. Hb4-2b (93.6 mg) was subjected to ODS C. C. using a MeOH– H_2O mixture to give 3 fractions (Hb4-2b-1–Hb4-2b-3). Fr. Hb4-2b-2 (48.3 mg) was further separated by Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to afford chichipenoside D (**1**, 16.7 mg, 0.003%, Fr. Hb4-2b-2d).

Fr. Hb4-2d (1.0 g) was subjected to ODS C. C. using a MeOH– H_2O mixture to give 3 fractions (Hb4-2d-1–Hb4-2d-3). Fr. Hb4-2d-2 (905.5 mg) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to give 3 fractions (Hb4-2d-2a–Hb4-2d-2c). Fr. Hb4-2d-2b (202.3 mg) was further separated by ODS C. C. using a MeCN– H_2O mixture to afford oleanolic acid 3-*O*- β -D-xylopyranosyl-(1 \rightarrow 2)- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranosyl-28-*O*- β -D-glucopyranoside (**6**, 72.0 mg, 0.011%, Fr. Hb4-2d-2b-9).

Fr. Hb2 (853.5 mg) was subjected to ODS C. C. using a MeOH– H_2O mixture to afford 5 fractions (Hb2-1–Hb2-5) and oleanolic acid 3-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 3)- β -D-glucuronopyranosyl-28-*O*- β -D-glucopyranoside (**8**, 440.3 mg, 0.069%, Fr. Hb2-4).

Fr. Hb2-2 (86.5 mg) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to afford cochalinoside C (**5**, 27.5 mg, 0.004%, Fr. Hb2-2e).

The 100% MeOH-eluted fraction (94.5 g) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to give 8 fractions (1–8). Fr. 4 (1.9 g) was subjected to ODS C. C. using a MeCN– H_2O mixture to give 3 fractions (4A–4C). Fr. 4B (1.2 g) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to afford oleanolic acid 3-*O*- α -L-rhamnopyranosyl-(1 \rightarrow 3)-6'-*O*-methyl- β -D-

glucuronopyranosyl-28-*O*- β -D-glucopyranoside (**9**, 40.5 mg, 0.006%, Fr. 4B-2).

Fr. 5 (15.9 g) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to give 6 fractions (5A–5F). Fr. 5E (300.8 mg) was subjected to ODS C. C. using a MeOH– H_2O mixture to give 2 fractions (Fr. 5E-1–5E-2). Fr. 5E-1 (235.2 mg) were partially (46.0 mg) subjected to ODS C. C. using a MeCN– H_2O mixture to afford longispinoside A (**2**, 16.0 mg, 0.003%, Fr. 5E-1e).

Fr. 5B (186.1 mg) was subjected to ODS C. C. using a MeCN– H_2O mixture to give 3 fractions (5B-1–5B-3). Fr. 5B-2 (51.4 mg) was subjected to ODS C. C. using a MeOH– H_2O mixture to afford longispinoside A methyl ester (**3**, 24.2 mg, 0.004%, Fr. 5B-2c).

Fr. 5D (391.3 mg) was subjected to ODS C. C. using a MeOH– H_2O mixture to give 3 fractions (5D-1–5D-3). Fr. 5D-2 (54.3 mg) was subjected to ODS C. C. using a MeOH– H_2O mixture to give 2 fractions (5D-2a–5D-2b). Fr. 5D-2b (40.3 mg) was subjected to ODS C. C. using a MeCN– H_2O mixture to afford erythronoside A methyl ester (**4**, 9.3 mg, 0.001%, Fr. 5D-2b-6).

Fr. 8 (6.1 g) was subjected to Si. C. C. using a CHCl_3 –MeOH– H_2O mixture to give 2 fractions (8A–8B). Fr. 8B (1.6 g) was subjected to ODS C. C. using a MeOH– H_2O mixture to afford stenoside A (**7**, 28.9 mg, 0.005%, Fr. 8B-3).

Chichipenoside D (**1**): colorless amorphous powder; $[\alpha]_{\text{D}}^{21}$ -28.0 (*c* 0.25, MeOH); IR (ATR) ν_{max} : 3385, 2939, 1714, 1602, 1417, 1376, 1293, 1041 cm^{-1} ; ^1H - and ^{13}C -NMR (DMSO- d_6 , 500 and 125 MHz), see Tables 2 and 3; negative HRFABMS *m/z*: 1103.5640 $[\text{M}-\text{H}]^-$ (calcd for $\text{C}_{54}\text{H}_{87}\text{O}_{23}$, 1103.5640).

Longispinoside A (**2**): colorless amorphous powder, $[\alpha]_{\text{D}}^{18}$ -52.0 (*c* 0.2 MeOH); IR (ATR) ν_{max} : 3330, 2940, 1740, 1450, 1020 cm^{-1} ; ^1H - and ^{13}C -NMR (DMSO- d_6 , 500 and 125 MHz), see Tables 2 and 3; positive HRFABMS *m/z*: 803.4546 $[\text{M}+\text{Na}]^+$ (calcd for $\text{C}_{42}\text{H}_{68}\text{O}_{13}\text{Na}$, 803.4548).

Longispinoside A methyl ester (**3**): colorless amorphous powder, $[\alpha]_{\text{D}}^{19}$ -117.3 (*c* 0.1 MeOH); IR (ATR) ν_{max} : 3363, 2945, 1746, 1442, 1024 cm^{-1} ; ^1H - and ^{13}C -NMR (DMSO- d_6 , 500 and 125 MHz), see Tables 2 and 3; positive HRFABMS *m/z*: 795.4900 $[\text{M}+\text{H}]^+$ (calcd for $\text{C}_{43}\text{H}_{71}\text{O}_{13}$, 795.4895).

Erythronoside A methyl ester (**4**): colorless amorphous powder, $[\alpha]_{\text{D}}^{19}$ -4.1 (*c* 0.1 MeOH); IR (ATR) ν_{max} : 3330, 2970, 1752, 1040 cm^{-1} ; ^1H - and ^{13}C -NMR (DMSO- d_6 , 500 and 125 MHz), see Tables 2 and 3; positive HRFABMS *m/z*: 941.5481 $[\text{M}+\text{H}]^+$ (calcd for $\text{C}_{43}\text{H}_{71}\text{O}_{13}$, 941.5474).

Cochalinoside C (**5**): colorless amorphous powder, $[\alpha]_{\text{D}}^{22}$ -28.8 (*c* 0.25 MeOH); IR (ATR) ν_{max} : 3370, 2970, 2940, 1710, 1610, 1420, 1390, 1240, 1140, 1050 cm^{-1} ; ^1H - and ^{13}C -NMR (DMSO- d_6 , 500 and 125 MHz), see Tables 2 and 3; negative HRFABMS *m/z*: 939.4952 $[\text{M}-\text{H}]^-$ (calcd for $\text{C}_{48}\text{H}_{75}\text{O}_{18}$, 939.4962).

Table 2 ¹H-NMR spectroscopic data for compounds 1–5,7 (500 MHz, in DMSO-d₆)

Position	1	2	3	4	5	7
1	α β	0.88 (o) 1.49 (o)	0.87 (o) 1.47 (o)	0.88 (o) 1.49 (o)	0.84 (o) 1.49 (o)	α β
2	α β	1.88 (brd, 10.8) 1.49 (o)	1.73 (brd, 10.2) 1.52 (o)	1.62 (o) 1.52 (o)	1.68 (o) 1.53 (o)	α β
3		3.02 (dd, 11.3, 3.2)	3.05 (dd, 11.3, 4.0)	3.05 (dd, 11.5, 4.4)	3.04 (dd, 11.3, 4.0)	3.03 (dd, 9.9, 3.8)
5		0.71 (brd, 11.3)	0.71 (brd, 11.6)	0.71 (brd, 11.6)	0.69 (brd, 11.5)	0.71 (o)
6	α β	1.51 (o) 1.35 (o)	1.49 (o) 1.33 (o)	1.49 (o) 1.32 (o)	1.47 (o) 1.33 (brt, 11.5)	α β
7	α β	1.48 (o) 1.24 (o)	1.47 (o) 1.25 (o)	1.47 (o) 1.26 (o)	1.47 (o) 1.26 (o)	α β
9		1.47 (o)	1.45 (o)	1.45 (o)	1.49 (o)	1.43 (o)
11		1.81 (brs)	1.78 (o)	1.78 (o)	1.78 (o)	1.74 (o)
12		5.21 (brs)	5.14 (t-like, 3.2)	5.14 (t-like, 3.4)	5.10 (t-like, 3.3)	5.20 (brs)
15	α β	1.12 (o) 1.47 (o)	1.20 (dd, 12.4, 4.4) 1.61 (t, 12.4)	1.20 (dd, 13.0, 4.8) 1.61 (t, 13.0)	0.84 (o) 1.69 (o)	α β
16		4.42 (m)	4.06 (dd, 12.4, 4.4)	4.06 (m)	1.72 (o)	4.06 (t-like, 12.9)
16-OH				4.38 (d, 6.2)		
18	α	2.50 (m)	2.21 (dd, 13.3, 4.3)	2.22 (dd, 12.0, 3.5)	1.92 (dd, 13.1, 3.8)	2.89 (dd, 13.4, 3.8)
19	β	1.68 (t, 13.1)	1.61 (t, 13.3)	1.61 (o)	1.66 (t, 13.1)	1.63 (t, 13.4)
21	α β	0.93 (o) 1.56 (brt, 12.6)	0.96 (o) 1.30 (o)	0.98 (o) 1.35 (o)	0.96 (o) 1.22 (o)	β α
22	β	1.68 (brd, 12.6)	1.08 (brd, 9.1)	1.08 (o)	1.07 (o)	β
23	α	4.25 (dd, 12.6, 4.5)	1.88 (brd, 13.5)	1.88 (dt, 13.6, 3.0)	1.44 (o)	α
24	β	0.96 (s)	1.37 (td, 13.5, 3.4)	1.38 (td, 13.6, 3.4)	1.25 (o)	β
25		0.76 (s)	0.96 (s)	0.96 (s)	0.95 (s)	0.95 (s)
26		0.88 (s)	0.87 (s)	0.87 (s)	0.74 (s)	0.74 (s)
27		0.90 (s)	0.90 (s)	0.90 (s)	0.87 (s)	0.84 (s)
28		1.16 (s)	1.14 (s)	1.14 (s)	1.10 (s)	1.13 (s)
28-OH		3.33 (o)	3.12 (o)	3.13 (o)	2.93 (o)	
29		3.77 (m)	3.51 (d, 10.2)	3.51 (dd, 10.5, 4.0)	3.29 (dd, 9.6, 4.8)	
30		0.88 (s)	3.46 (o)	3.46 (o)	4.28 (t, 4.8)	0.90 (s)
1'		0.87 (s)	0.84 (s)	0.85 (s)	0.84 (s)	0.86 (s)
		GlcA	GlcA	GlcA (Me)	GlcA (Me)	0.85 (s)
		4.17 (d, 7.7)	4.25 (d, 7.7)	4.31 (d, 7.9)	4.52 (d, 8.3)	GlcA
					4.16 (d, 7.6)	4.30 (d, 7.4)

Table 2 (continued)

Position	1	2	3	4	5	7
2'	3.08 (o)		3.11 (o)	3.12 (o)	3.66 (t, 8.3)	3.08 (q, 7.6)
2'-OH						3.58 (o)
3'	3.36 (o)		3.38 (t, 9.3)	3.40 (t, 8.9)	3.58 (t, 8.3)	5.17 (d, 7.6)
4'	3.17 (o)		3.29 (t, 9.3)	3.32 (o)	3.51 (td, 8.3, 6.2)	3.36 (o)
4'-OH					5.22 (d, 6.2)	3.19 (t, 7.6)
5'	3.31 (o)		3.57 (d, 9.3)	3.80 (d, 9.4)	3.88 (d, 8.3)	3.33 (d, 7.6)
6'-OCH ₃				3.65 (s)		3.35 (o)
1''	Rha-1		Rha	Rha	Glc	Glc-1
2''	5.04 (brs)		5.02 (brs)	5.00 (d, 1.4)	4.43 (d, 7.9)	4.46 (d, 9.1)
2''-OH			3.69 (brs)	3.69 (td, 4.0, 1.4)	2.98 (o)	2.95 (o)
3''	3.45 (o)		3.47 (dd, 9.4, 3.1)	4.55 (d, 4.0)		4.86 (d, 4.3)
3''-OH				3.46 (m)	3.13 (t, 8.9)	3.13 (o)
4''	3.15 (o)		3.16 (t, 9.4)	3.16 (td, 9.6, 4.7)	2.96 (o)	5.05 (brs)
4''-OH				4.55 (d, 4.7)		2.96 (o)
5''	3.96 (dq, 9.4, 6.1)		3.88 (dq, 9.4, 6.2)	3.85 (dq, 9.6, 6.2)	3.03 (o)	4.96 (d, 4.5)
6''	1.05 (d, 6.1)		1.06 (d, 6.2)	1.06 (d, 6.2)	3.39 (m)	3.03 (o)
6''-OH					3.66 (o)	3.38 (o)
1'''	Glc				Rha	3.65 (dd, 9.4, 5.4)
2'''	4.33 (d, 7.8)				4.94 (brs)	5.01 (o)
2'''-OH					3.75 (t-like, 3.3)	Rha
3'''	5.26 (d, 3.3)					5.74 (d, 1.1)
4'''	3.21 (o)					3.55 (brs)
4'''-OH	3.36 (o)				3.45 (brd, 10.2)	3.21 (o)
5'''	3.11 (m)				3.18 (td, 10.2, 4.5)	3.21 (o)
6'''	3.45 (o)				4.61 (d, 4.5)	3.44 (o)
6'''-OH	3.57 (brd, 11.6)				3.80 (dq, 10.2, 6.2)	3.16 (o)
1''''	Rha-2				1.07 (d, 6.2)	3.50 (o)
2''''	4.69 (brs)					1.08 (d, 6.2)
3''''	3.59 (brs)					Glc-2
4''''	3.39 (o)					5.32 (d, 7.9)
4''''-OH	3.17 (o)					3.40 (o)
						3.45 (o)
						3.16 (o)
						5.14 (o)

Table 2 (continued)

Position	1	2	3	4	5	7
5 ^{'''}		3.83 (dq, 9.2, 6.1)				3.18 (o)
6 ^{'''}		1.08 (d, 6.1)				3.42 (o)
						3.59 (o)
1 ^{''''}						Xyl
2 ^{''''}						4.48 (d, 7.9)
2 ^{''''} -OH						2.93 (o)
3 ^{''''}						5.30 (d, 4.0)
3 ^{''''} -OH						3.07 (td, 9.5, 3.9)
4 ^{''''}						5.08 (d, 3.9)
5 ^{''''}						3.18 (o)
						2.98 (o)
						α
						β

Oleanolic acid 3-*O*-β-D-xylopyranosyl-(1 → 2)-α-L-rhamnopyranosyl-(1 → 3)-β-D-glucuronopyranosyl-28-*O*-β-D-glucopyranoside (**6**): colorless amorphous powder, $[\alpha]_{\text{D}}^{23}$ -4.9 (*c* 0.3 MeOH); IR (ATR) ν_{max} : 3370, 2970, 2930, 1730, 1460, 1390, 1260, 1230, 1020 cm^{-1} ; ^1H - and ^{13}C -NMR (DMSO-*d*₆, 500 and 125 MHz), see Tables 2 and 3; negative HRFABMS *m/z*: 1071.5386 $[\text{M}-\text{H}]^-$ (calcd for C₅₃H₈₃O₂₂, 1071.5309).

Stenoside A (**7**): colorless amorphous powder, $[\alpha]_{\text{D}}^{19}$ -8.6 (*c* 0.1 MeOH); IR (ATR) ν_{max} : 3372, 2927, 1732, 1602, 1417, 1260, 1074, 1041 cm^{-1} ; ^1H - and ^{13}C -NMR (DMSO-*d*₆, 500 and 125 MHz), see Tables 2 and 3; positive HRFABMS *m/z*: 1273.5642979.4658 $[\text{M}+\text{K}]^+$ (calcd for C₅₉H₉₄O₂₇K, 1273.5620).

Alkaline hydrolysis of 8

Oleanolic acid 3-*O*-α-L-rhamnopyranosyl-(1 → 3)-β-D-glucuronopyranosyl-28-*O*-β-D-glucopyranoside (**8**) (10.1 mg) was dissolved in 4 mL of MeOH and hydrolyzed with 0.4 mL of 5% K₂CO₃ aq. solution under reflux for 4 h at 110 °C. After the reaction, the mixture was subjected to DOWEX 50 W-X8 to remove potassium ions. After washing with plenty of MeOH, the eluent was evaporated to dryness. The residue was chromatographed over ODS C. C. using a stepwise gradient (80% MeOH → 100% MeOH) to afford hydrolysate (**8a**) (2.0 mg), the 28-*O*-deglycosylated saponin of **8**.

Determination of sugar configuration

Sugar configuration was determined according to a previously reported procedure [25]. Saponins (**1–7**) (each 2 mg) were dissolved in 1 M HCl (0.4 mL) and then heated at 110 °C for 2 h. Following neutralization with 1 M NaOH, the reaction mixture was evaporated under reduced pressure to dryness, and then the residue was dissolved in pyridine (0.4 mL) containing L-cysteine methyl ester hydrochloride (2 mg) and heated at 60 °C for 1 h. *O*-Tolyliisothiocyanate (2 μL) was then added to the mixture, and it was heated at 60 °C for 1 h. The reaction mixture was analyzed by HPLC using a system equipped with a PU-2089 pump, UV-2075 UV/VIS detector (Nippon Bunko Co.), and Senshu Pak PEGASIL 4.6 φ × 250-mm HPLC column (temp, 35 °C; flow, 0.8 mL/min; elution buffer, MeCN–H₂O 25:75 containing 50 mM H₃PO₄). The HPLC column was washed with MeOH after each injection. The reaction conditions for D, L-glucose, D-glucuronic acid, D, L-xylose, and L-rhamnose were the same as described above except for the mass, authentic sugar (5 mg), and L-cysteine methyl ester hydrochloride (5 mg), respectively. L-Glucuronic acid and D-rhamnose derivatives were synthesized from D-glucuronic acid,

Table 3 ^{13}C -NMR spectroscopic data for compounds **1–5,7** (125 MHz in $\text{DMSO-}d_6$)

Position	1	2	3	4	5	7						
1	38.3	CH ₂	38.2	CH ₂	38.1	CH ₂	38.1	CH ₂	38.2	CH ₂	38.3	CH ₂
2	25.4	CH ₂	25.6	CH ₂	25.6	CH ₂	25.4	CH ₂	25.4	CH ₂	25.4	CH ₂
3	88.1	CH	88.3	CH	88.4	CH	88.6	CH	88.1	CH	88.7	CH
4	38.7	C	38.7	C	38.7	C	38.6	C	38.7	C	38.7	C
5	54.8	C	54.8	CH	54.8	CH	54.8	CH	55.0	CH	55.2	CH
6	17.7	CH ₂	17.8	CH ₂	17.7	CH ₂	17.8	CH ₂	17.7	CH ₂	17.9	CH ₂
7	32.1	CH ₂	32.2	CH ₂	32.1	CH ₂	32.1	CH ₂	32.4	CH ₂	32.3	CH ₂
8	39.4	C	39.2	C	39.1	C	39.9	C	39.2	C	39.1	C
9	46.2	CH	46.2	CH	46.2	CH	47.0	CH	46.2	CH	47.1	CH
10	36.1	C	36.1	C	36.1	C	36.1	C	36.2	C	36.3	C
11	23.1	CH ₂	23.0	CH ₂	23.0	CH ₂	23.0	CH ₂	23.0	CH ₂	23.0	CH ₂
12	122.4	CH	121.4	CH	121.6	CH	121.5	CH	122.2	CH	121.7	CH
13	141.9	C	143.3	C	143.3	C	144.3	C	142.4	C	143.5	C
14	41.7	C	42.9	C	42.9	C	41.2	C	43.3	C	41.2	C
15	34.8	CH ₂	35.2	CH ₂	35.2	CH ₂	25.1	CH ₂	36.5	CH ₂	27.6	CH ₂
16	64.5	CH	64.5	CH	64.5	CH	21.7	CH ₂	62.7	CH	22.1	CH ₂
17	44.6	C	40.1	C	40.1	C	36.5	C	50.1	C	46.0	C
18	42.0	CH	42.6	CH	42.6	CH	41.8	CH	43.3	CH	40.8	CH
19	45.5	CH ₂	46.4	CH ₂	46.3	CH ₂	46.3	CH ₂	45.1	CH ₂	45.5	CH ₂
20	31.4	C	30.5	C	30.5	C	30.7	C	30.2	C	30.3	C
21	41.0	CH ₂	33.5	CH ₂	33.5	CH ₂	33.8	CH ₂	32.4	CH ₂	33.3	CH ₂
22	77.9	CH	23.8	CH ₂	23.8	CH ₂	30.9	CH ₂	25.9	CH ₂	31.5	CH ₂
23	27.6	CH ₃	27.5	CH ₃	27.4	CH ₃	27.5	CH ₃	27.5	CH ₃	27.6	CH ₃
24	16.4	CH ₃	16.3	CH ₃	16.3	CH ₃	16.1	CH ₃	16.4	CH ₃	16.2	CH ₃
25	15.3	CH ₃	15.3	CH ₃	15.3	CH ₃	15.3	CH ₃	15.1	CH ₃	15.2	CH ₃
26	16.6	CH ₃	16.5	CH ₃	16.5	CH ₃	16.4	CH ₃	16.8	CH ₃	16.7	CH ₃
27	26.8	CH ₃	26.6	CH ₃	26.6	CH ₃	25.6	CH ₃	26.7	CH ₃	25.6	CH ₃
28	56.7	CH ₂	65.2	CH ₂	65.0	CH ₂	67.4	CH ₂	174.5	C	175.3	C
29	32.8	CH ₃	33.1	CH ₃	33.1	CH ₃	33.1	CH ₃	32.7	CH ₃	32.8	CH ₃
30	24.4	CH ₃	23.7	CH ₃	23.7	CH ₃	23.5	CH ₃	23.7	CH ₃	23.4	CH ₃
	GlcA		GlcA		GlcA (Me)		GlcA (Me)		GlcA		GlcA	
1'	105.1	CH	105.2	CH	105.2	CH	103.1	CH	105.2	CH	103.6	CH
2'	74.6	CH	74.3	CH	74.2	CH	77.0	CH	74.5	CH	77.7	CH
3'	80.3	CH	80.4	CH	80.1	CH	81.1	CH	80.3	CH	81.6	CH
4'	70.6	CH	70.2	CH	70.1	CH	69.9	CH	70.3	CH	70.9	CH
5'	74.2	CH	75.1	CH	75.2	CH	74.8	CH	74.1	CH	74.0	CH
6'	173.1	C	171.3	C	169.5	C	169.5	C	173.5	C	173.7	C
6'-OCH ₃					51.9	CH ₃	51.9	CH ₃				
	Rha-1		Rha		Rha		Glc		Rha-1		Glc-1	
1''	100.2	CH	100.5	CH	100.7	CH	101.6	CH	100.3	CH	101.6	CH
2''	70.6	CH	70.6	CH	70.5	CH	70.4	CH	70.6*	CH	70.4	CH
3''	70.7	CH	70.6	CH	70.5	CH	76.6	CH	70.6	CH	76.6	CH
4''	72.1	CH	72.1	CH	72.0	CH	74.2	CH	72.1	CH	74.6	CH
5''	67.8	CH	68.0	CH	68.1	CH	77.0	CH	67.8	CH	77.0	CH
6''	17.9	CH ₃	17.9	CH ₃	17.8	CH ₃	61.4	CH ₂	17.9	CH ₃	61.4	CH ₂
	Glc						Rha		Rha-2		Rha	
1'''	103.3	CH					100.5	CH	93.3	CH	100.6	CH
2'''	74.1	CH					70.6	CH	69.6	CH	70.6	CH
3'''	75.3	CH					70.6	CH	71.5	CH	70.8	CH
4'''	76.6	CH					71.9	CH	71.3	CH	72.1	CH
5'''	75.5	CH					68.6	CH	70.7	CH	68.2	CH

Table 3 (continued)

Position	1	2	3	4	5	7			
6'''	60.1	CH ₂		17.8	CH ₃	17.9	CH ₃	17.9	CH ₃
		Rha-2						Glc-2	
1''''	100.5	CH						91.9	CH
2''''	70.7	CH						78.8	CH
3''''	70.6	CH						76.9	CH
4''''	71.9	CH						77.6	CH
5''''	68.6	CH						69.2	CH
6''''	17.7	CH ₃						60.6	CH ₂
								Xyl	
1''''								104.0	CH
2''''								74.2	CH
3''''								76.5	CH
4''''								69.5	CH
5''''								65.8	CH ₂

L-rhamnose (each 5 mg), and D-cysteine methyl ester hydrochloride (5 mg). These derivatives were used for determining the retention time. The retention times (min.) for the authentic sugar derivatives were D-glucose (23.56), L-glucose (21.89), D-glucuronic acid (24.37), L-glucuronic acid (derived from D-glucuronic acid and D-cysteine methyl ester, 23.38), D-rhamnose (derived from L-rhamnose and D-cysteine methylester, 18.36), L-rhamnose (40.19), D-xylose (22.17), and L-xylose (20.14). These retention times were compared with the retention times of each saponin (1–7) in the reaction mixtures. The peaks at 23.55, 24.25, and 40.18 min of the sugar derivatives from **1** coincided with the derivatives of D-glucose, D-glucuronic acid, and L-rhamnose, respectively. The other saponins (2–7) provided the same results as **1**.

Th-T assay

The rate of A β aggregation was evaluated using a slightly modified thioflavin-T (Th-T) method described in a previous report [6]. The Th-T method was originally developed by Naiki and co-workers [48]. The rate of A β aggregation was calculated by comparing the fluorescence intensity of each sample with that of a control (25 μ M of A β ₄₀ and DMSO containing no test sample). The aggregation rate (%) was calculated using the following formula: $[(S - B)/(C - B) \times 100]$, (S, fluorescence of Th-T solution incubated with A β ₄₀ and sample; C, fluorescence of Th-T solution incubated with A β ₄₀ and DMSO; B, fluorescence of Th-T solution not incubated with A β ₄₀ and DMSO). Myricetin (Fujifilm-Wako) was used as a positive control at 25 μ M as the final concentration, and that of test samples (1–4, 7–8) was 100 μ M.

BACE1 FRET assay

BACE1 assays were performed in 384-well black plates using a BACE1 FRET assay kit, Red (Thermo Fisher Scientific, USA). The assay was carried out according to the supplied manual, with some modifications. Samples were dissolved in the assay buffer (50 mM sodium acetate, pH 4.5) with DMSO (final concentrations were 10%). Next, 9 μ L of test sample, 9 μ L of BACE1 substrate (750 nM Rh-EVNLDAEFK-Quencher, in 50 mM ammonium bicarbonate), and 9 μ L of BACE1 enzyme (1.0 U/mL) were mixed in the wells and incubated 60 min in the dark at room temperature. The fluorescence intensity of each well was measured using a SYNERGY HTX multimode reader (BioTek, USA), with excitation at 540 nm and emission at 590 nm. The inhibition ratio was calculated by the following formula: $\text{Inhibition rate (\%)} = \{1 - [(S - S_0) - (B - B_0)] / [(C - C_0) - (B - B_0)]\} \times 100$, where C represents the fluorescence of a control [enzyme, substrate, and assay buffer concentration with DMSO (final concentrations 10%)] after 60 min of incubation, C₀ represents the initial fluorescence of a control [enzyme, substrate, and assay buffer concentration with DMSO (final concentrations 10%)], B represents the fluorescence of a control [substrate and assay buffer concentration with DMSO (final concentrations 10%)] after 60 min of incubation, B₀ represents the initial fluorescence of a control [substrate and assay buffer concentration with DMSO (final concentrations were 10%)], S represents the fluorescence of the tested sample (enzyme, sample solution, and substrate) after 60 min of incubation, and S₀ represents the initial fluorescence of the tested sample (enzyme, sample solution, and substrate). β -Secretase inhibitor IV (Merck, Germany) was used as a positive

control at 100 μM . Isolated compounds (**1–4,7–8**) were tested for their BACE1 inhibitory activity at 100 μM (final concentration).

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