Four New Flavonoids Isolated from the Aerial Parts of Cadaba rotundifolia Forssk. (Qadab)

Gadah Abdulaziz Al-Hamoud 1,2, Raha Saud Orfali 2,* , Sachiko Sugimoto 1, Yoshi Yamano 1, Nafee Alothyqi 3, Ali Mohammed Alzahrani 4 and Katsuyoshi Matsunami 1,*

1 Department of Pharmacognosy, Graduate School of Biomedical and Health Sciences, Hiroshima University, 1-2-3 Kasumi, Minami-ku, Hiroshima 734-8553, Japan; galhamoud@ksu.edu.sa (G.A.A.-H.); ssugimoto@hiroshima-u.ac.jp (S.S.); yamano@hiroshima-u.ac.jp (Y.Y.)
2 Department of Pharmacognosy, College of Pharmacy, King Saud University, 11495 Riyadh, Saudi Arabia
3 Department of Biology, Umm Al-Qura University, 1109 Makkah Al-Mukarramah, Saudi Arabia; niothyqi@uqu.edu.sa
4 Department of Biology, Arts and Sciences in Qilwah, Al-Baha University, 1988 Al-Baha, Saudi Arabia; alialzahrani@bu.edu.sa
* Correspondence: rorfali@ksu.edu.sa (R.S.O.); matunami@hiroshima-u.ac.jp (K.M.); Tel: +966-11-8055014 (R.S.O.); +81-82-257-5335 (K.M.)

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Abstract: Cadaba rotundifolia (Forssk.) (family: Capparaceae; common name: Qadab) is one of four species that grow in the Red Sea coastal region in the Kingdom of Saudi Arabia. The roots and leaves of C. rotundifolia is traditionally used to treat tumors and abscesses in Sudan. A previous phytochemical study of the roots yielded a quaternary alkaloid, but no report on chemical constituents of the aerial parts of C. rotundifolia growing in Saudi Arabia has been issued so far. Oxidative stress and advanced glycation end products (AGEs) are thought as causal factors in many degenerative diseases, such as Alzheimer’s disease, diabetes, atherosclerosis and aging. In this study, a total of twenty compounds, including four previously undescribed acylated kaempferol glucosides, were isolated from the aerial parts of C. rotundifolia collected in Saudi Arabia. These new compounds were identified as kaempferol 3-O-[2-(trans-feruloyl)-3-O-β-d-glucopyranosyl]-β-d-glucopyranoside (1), kaempferol 3-O-β-neohesperidoside-7-O-[2-O-(cis-p-coumaroyl)-3-O-β-d-glucopyranosyl]-β-d-glucopyranoside (2), kaempferol 3-O-[2,6-di-O-α-l-rhamnopyranosyl]-β-d-glucopyranoside-7-O-[6-O-(trans-feruloyl)]-β-d-glucopyranoside (3) and kaempferol 3-O-[2,6-di-O-α-l-rhamnopyranosyl]-β-d-glucopyranoside-7-O-[6-O-(trans-p-coumaroyl)]-β-d-glucopyranoside (4). Their structures were established based on UV-visible, 1D, 2D NMR, and HR-ESI-MS analyses. Of the assayed compounds, 17 and 18 showed potent radical scavenging activity with IC_{50} values of 14.5 and 11.7 µM, respectively, and inhibitory activity toward AGEs together with compound 7 with IC_{50} values 96.5, 34.9 and 85.5 µM, respectively.

Keywords: Cadaba rotundifolia; Capparaceae; Qadab; aerial parts; flavonoid; antioxidant; AGEs

1. Introduction

Oxidative stress and advanced glycation end products (AGEs) are thought as causal factors of many diseases, such as diabetes, atherosclerosis, neurodegenerative disorders and aging. Therefore, the discovery of antioxidant and anti-AGEs molecules are important for preventing these age-related pathogenesis. Cadaba (Forssk.) is one of genera in the Capparaceae family, with about 30 species distributed in southern Africa, India, Malaysia, and Australia [1,2]. Among them four species grow in the Red Sea coastal region in the Kingdom of Saudi Arabia [3]. Phytochemical studies of several Cadaba species led to isolation of various secondary metabolites, such as alkaloids, terpenoids...
and flavonoids [4–7]. *Cadaba rotundifolia* Forssk., known by the common name Qadab, is an erect shrub with numerous drooping branches with young twigs covered with short glandular hair [8]. In Sudan, *C. rotundifolia* roots and leaves are traditionally used to treat tumors and abscesses. A previous chemical study of the ethanolic extract of *C. rotundifolia* root yielded a quaternary alkaloid (3-hydroxystachydrine) [9]. However, no report on chemical constituents of the aerial parts of *C. rotundifolia* has been issued so far. In this work, we have investigated the chemical profile of the aerial parts of *C. rotundifolia* collected in Saudi Arabia. As a result, twenty flavonoids were obtained, including four undescribed acylated kaempferol glucosides 1–4 and 16 known compounds. This paper describes on the isolation, structural elucidation, and evaluation of antioxidant (DPPH) and inhibitory activity toward AGEs formation.

2. Results and Discussion

2.1. Isolation and Spectroscopic Analyses of the Compounds

The aerial parts of *C. rotundifolia* were extracted with MeOH at room temperature till exhausted. The MeOH extract (800 g) was further separated by solvent fractionation, various column chromatographies (CC), and HPLC to afford compounds 1–20 (Figure 1). The chemical structures of the isolated compounds were investigated and identified through intensive spectroscopic analyses based on UV-visible, 1D and 2D NMR, and HR-ESI-MS data as follows (Figures S1–S35).

![Chemical structures of compounds isolated from *Cadaba rotundifolia*.](image)

**Figure 1.** Compounds 1–20 isolated from *Cadaba rotundifolia*.

2.1.1. Chemical Structure of Compound 1

Compound 1 was isolated as a yellow amorphous powder, soluble in MeOH with a negative optical rotation ([α]D25 −79.0). Its molecular formula, C37H38O19Na, was determined by positive HR-ESI-MS analysis (m/z 809.1888 [M + Na]+, calcd for C37H38O19Na 809.1899). It had a UV spectrum typical
of an acylated kaempferol glycoside ($	ext{λ}_{\text{max}}$ 327, 267 nm) [10,11]. The IR spectrum suggested the presence of hydroxy (3388 cm$^{-1}$), α,β-unsaturated carbonyl ester (1710 cm$^{-1}$), α,β-unsaturated ketone (1658 cm$^{-1}$), aromatic ring (1513, 1446 cm$^{-1}$) and ether (1176, 1070 cm$^{-1}$) functions. The proton and carbon resonances in its $^1$H and $^{13}$C-NMR spectra clarified the aromatic and glycosidic nature of 1. It was found that the $^1$H-NMR spectrum of 1 displayed feature resembling to those of 5 [12], as both the aglycone and glycoside units were almost identical. The major differences were in the signals related to the acyl group of 5. The $^1$H-NMR spectrum of 1 (Table 1) exhibited meta-coupled resonances at $\delta_{\text{H}}$ 6.17 and 6.37 (each 1H, d, $J = 2.0$ Hz), characteristic of H-6 and H-8 of the A-ring, respectively. The $^1$H-$^1$H correlation spectrum (COSY, Figure 2) and $^1$H-NMR spectrum of 1 displayed two ortho-coupled doublet resonances at $\delta_{\text{H}}$ 8.01 and 6.91 (each 2H, d, $J = 8.8$ Hz) assignable to H-2′/6′ and H-3′/5′ of the B-ring, respectively, which indicated the presence of a kaempferol aglycone (Table 1).

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m: multiplet or overlapped signals.

The $^1$H-NMR spectrum of 1 (Table 1) also gave signals at $\delta_{\text{H}}$ 6.83 (1H, d, $J = 8.2$ Hz), 7.09 (1H, d, $J = 8.2$, 1.8 Hz) and 7.20 (1H, d, $J = 1.8$ Hz) attributable to an ABX coupling system for a 1,3,4-trisubstituted aromatic moiety, and two trans-coupled olefinic protons at $\delta_{\text{H}}$ 6.40 (1H, br d, $J = 15.9$ Hz) and 7.68 (1H, br d, $J = 15.9$ Hz), together with a singlet signal for a methoxy group at $\delta_{\text{H}}$ 3.92 which suggested the presence of a feruloyl or isoferuloyl function. The two anomic proton resonances, which occur at $\delta_{\text{H}}$ 5.73 (1H, d, $J = 8.0$ Hz) and 4.43 (1H, d, $J = 7.8$ Hz) and 12 carbon signals (δc 104.9, 100.5, 84.9, 78.4, 78.1, 77.7, 74.8, 74.6, 71.4, 70.0, 62.5 and 62.4) correspond to two glucose
moeities. Acid hydrolysis of 1 released d-glucose and the β-anomeric configuration for the sugars was determined from the magnitude of the $J_{1,2}$ coupling constants. In addition, alkaline hydrolysis afforded ferulic acid, which was identified by comparing with an authentic sample based on their HPLC retention times. The $^{13}C$-NMR spectrum (Table 2) of 1 showed 37 carbon resonances, 12 of which assigned to two β-glucose units, 15 signals were consistent with kaempferol aglycone and the remaining 10 signals were ascribed to acyl groups. The correlations observed between the proton and carbon resonances of the acyl moiety from H-7′′′′ ($\delta H$ 7.68) to C-2′′′′ ($\delta C$ 111.7), C-6′′′′ ($\delta C$ 124.3) and an ester carbonyl C-9′′′′ ($\delta C$ 168.5), and from the protons of methoxy group ($\delta H$ 3.92) to C-3′′′′ ($\delta C$ 149.3) in HMBC spectrum (Figure 2) indicated the presence of a feruloyl unit.

![Image](image-url)

**Figure 2.** COSY and HMBC correlations of 1–4.

According to the coupling constant of the olefinic protons at H-7′′′′ and H-8′′′′ ($J = 15.9$ Hz), the configuration of the feruloyl moiety was determined as trans. Full assignments of the $^1H$ and $^{13}C$ resonances of the structural fragments of 1 were achieved by analysis of COSY, HSQC and HMBC spectra. A correlation between the anemic H-1′′ proton ($\delta H$ 5.73) and $\delta C$ 134.8 in the HMBC spectrum (Figure 2) defined C-3 of kaempferol as a site of O-glucosylation. Moreover, the interglucosidic linkage of the disaccharide moiety was also determined by the HMBC cross-peak between H-3′′ ($\delta H$ 3.91) and C-1′′′′ ($\delta C$ 104.9). The E-feruloyl unit was linked to C-2′ based on the correlations of the H-2′′ ($\delta H$ 5.23) with C-1′′′ (δc 100.5), C-3′′ (δc 84.9) and ester carbonyl carbon C-9′′′′ (δc 168.5). Therefore, the structure of compound 1 was determined as kaempferol 3-O-[2-O-(trans-feruloyl)-3-O-β-d-glucopyranosyl]-β-d-glucopyranoside.
was similar to those of the kaempferol glycoside acylated with a hydroxycinnamic acid [10,11,13].

A-ring, respectively. In comparison with literature, the downfield shifts of H-6 and H-8 in the

1H-NMR spectra by four anomeric proton signals at

\[ \delta = 5.29 \text{ (1H, d, } J = 5.83 \text{ and } 6.86 \text{) with the ester carbonyl carbon at } \delta \text{. Similarly, the AA}

set of kaempferol signals, a -coupled proton resonances at

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Table 2. 13C-NMR data of compounds 1-4 (150 or 175 * MHz, CD3OD, } \delta \text{ in ppm).}

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2.1.2. Chemical Structure of Compound 2

Compound 2 was isolated as a yellow amorphous powder, with a positive optical rotation ([\alpha]^{22}_D +14.5). Its molecular formula was determined to be C_{48}H_{56}O_{27} by positive HR-ESI-MS (m/z = 1087.2875 ([M + Na]^{+}, calc for C_{48}H_{56}O_{27}Na 1087.2901). The UV profile of 2 displayed \lambda_{max} 316, 267 nm, which was similar to those of the kaempferol glycoside acylated with a hydroxycinnamic acid [10,11,13]. The IR spectrum indicated absorptions of hydroxyl (3397 cm\(^{-1}\)), \alpha,\beta-unsaturated carbonyl ester (1721 cm\(^{-1}\)), \alpha,\beta-unsaturated ketone (1657 cm\(^{-1}\)), aromatic ring (1512, 1442 cm\(^{-1}\)) and ether (1157, 1072 cm\(^{-1}\)) functions. The above features were confirmed by the \( ^1H \)-NMR spectrum which revealed a set of kaempferol signals, a p-coumaroyl group and glycoside moieties. The kaempferol aglycone of 2 was represented in the \( ^1H \)-NMR spectrum (Table 1) by two ortho-coupled resonances of the B-ring at \( \delta_H 8.06 \) and 6.89 (each 1H, d, \( J = 8.9 \) Hz), typical of H-2'/6' and H-3'/5', respectively, in addition to two meta-coupled proton resonances at \( \delta_H 6.37 \) and 6.65 (each 1H, br s), attributed to H-6 and H-8 of the A-ring, respectively. In comparison with literature, the downfield shifts of H-6 and H-8 in the \( ^1H \)-NMR spectrum suggested the presence of substitution at C-7 of kaempferol, while the substitution at C-3 was evident from the downfield shift of C-2 to \( \delta_C 159.2 \) in the \( ^1C \)-NMR spectrum. Moreover, the \( ^1H \)-NMR spectrum (Table 1) exhibited two olefinic protons at \( \delta_H 5.83 \) and 6.86 with a coupling constant of 12.9 Hz. Similarly, the AA'BB' aromatic coupling system at \( \delta_H 6.67 \) and 7.55 (each 1H, d, \( J = 8.6 \) Hz) were attributed to a p-coumaroyl unit, which was confirmed by the correlation of the olefinic protons (\( \delta_H 5.83 \) and 6.86) with the ester carbonyl carbon at \( \delta_C 167.2 \) and aromatic carbons (\( \delta_C 127.6; C-1'/**'**'**' and 133.4; C-2/6'/**'**'**'**'**' in the HMBC spectrum (Figure 2). The sugar units in 2 were represented in the \( ^1H \)-NMR spectra by four anomer proton signals at \( \delta_H 4.40 \) (1H, d, \( J = 7.8 \) Hz), 5.22 (1H, d, \( J = 1.3 \) Hz), 5.29 (1H, d, \( J = 7.8 \) Hz), 5.72 (1H, d, \( J = 7.6 \) Hz) and one methyl group at 0.96 (3H, d, \( J = 6.18 \) Hz),
which indicated the tetruglycosidic nature of 2. Identification of the sugar residues was carried out by careful analysis of the 1D and 2D NMR data as well as comparison with the reported chemical shift values. The absolute stereochemistry of the sugars was determined by acid hydrolysis of 2 and HPLC analysis with an optical rotation detector, and the magnitudes of \( f_{1,2} \) coupling constants that revealed \( \beta-L-glucopyranose \) and \( \alpha-L-rhamnopyranose \) moieties. The spectroscopic data of 2 (Tables 1 and 2) indicated that it is an isomer of 12 [13]. The relatively smaller coupling constant (\( f = 12.9 \) Hz) and the chemical shift values of H-7 \( \delta_1 \) and H-8 \( \delta_2 \) (6.86 and 5.83, respectively) indicated 2 as a cis isomer. In addition, alkaline hydrolysis of 2 afforded cis-p-coumaric acid. The positions of glycosylation and the cis-p-coumaroyl moiety on the kaempferol aglycone were decided using the HMBC spectrum (Figure 2). The anomic protons of two glucose moieties at \( \delta_1 \) 5.72 and 5.29 have long-range correlations with C-3 (\( \delta_1 \) 134.7) and C-7 (\( \delta_1 \) 163.8) of the kaempferol aglycone, respectively, indicating the connectivity of the sugars. The COSY correlation (Figure 2) of H-2* at (\( \delta_1 \) 3.60) with H-1* at (\( \delta_1 \) 5.72), together with correlation observed in the HMBC spectrum between H-2* and C-1* of rhamnose at (\( \delta_1 \) 102.6) confirmed the presence of a \( \beta-L-neohesperidoside \) structure at C-3 of kaempferol. Moreover, the correlations of H-2** (\( \delta_1 \) 5.25) with H-1** (\( \delta_1 \) 5.29) and H-3** (\( \delta_1 \) 3.92) in COSY spectrum, and the correlations from H-2’* to C-9’* (\( \delta_1 \) 167.2), and from H-3’* to C-1’* (\( \delta_1 \) 105.0) in HMBC spectrum confirmed the presence of 7-O-[2-O-(cis-p-coumaroyl)-3-O-\( \beta-L-glucopyranosyl]-\( \beta-L-glucopyranosyl \) as the kaempferol aglycone. The structure of 2 was therefore identified as kaempferol 3-O-\( \beta-L-neohesperidoside-7-O-[2-O-(cis-p-coumaroyl)-3-O-\( \beta-L-glucopyranosyl]-\( \beta-L-glucopyranoside \).}

### 2.1.3. Chemical Structure of Compound 3

Compound 3 was obtained as a yellow amorphous powder with a negative optical rotation (\( [\alpha]^{25}_D \) -82.5). The molecular formula, \( C_{49}H_{56}O_{27} \), was deduced from the sodiated molecular ion peak at \( m/z \) 1101.3021 (calcd for \( C_{49}H_{56}O_{27} \) 1101.3058) by HR-ESI-MS measurement. It displayed characteristic UV spectrum of an acylated kaempferol glycoside with \( \lambda_{max} \) at 329, 268 nm, similar to those of 1. The IR spectrum exhibited absorptions of hydroxy (3389 cm\(^{-1}\)), \( \alpha-\)unsaturated ketone (1651 cm\(^{-1}\)), aromatic ring (1513, 1455 cm\(^{-1}\)) and ether (1180, 1071 cm\(^{-1}\)) functions. Analysis of the \( ^1H \) and \( ^13C \)-NMR spectra of 3 (Tables 1 and 2) indicated the presence of a kaempferol residue, four hexose sugar moieties, and an E-feruloyl unit. The \( ^1H \) spectrum (Table 1) displayed four downfield doublet signals at \( \delta_1 \) 6.46 (1H, d, \( J = 2.1 \) Hz), 6.61 (1H, d, \( J = 2.1 \) Hz), 6.76 (2H, d, \( J = 8.9 \) Hz) and 7.93 (2H, d, \( J = 8.9 \) Hz) assignable to H-6, 8, 3,5/’ and 2,6/’, respectively, belonging to kaempferol aglycone. Moreover, the characteristic signals of the E-feruloyl moiety were observed in \( ^1H \)-NMR spectrum as five downfield doublets at \( \delta_1 \) 6.28 (1H, br d, \( J = 15.8 \) Hz), 6.67 (1H, d, \( J = 8.2 \) Hz), 6.85 (1H, d, \( J = 8.2 \) Hz), 6.98 (1H, d, \( J = 1.7 \) Hz) and 7.50 (1H, br d, \( J = 15.8 \) Hz), corresponding to H-8*, H-5**, H-6**, H-2’* and H-7**, respectively, in addition to a singlet signal of a methoxy group at \( \delta_1 \) 3.75. Full assignment of the 1D NMR spectra of 3 was achieved by analysis of COSY, HSQC and HMBC data, which confirmed presence of a 3,7-O-glycosidic kaempferol residue together with the E-feruloyl unit. Acid hydrolysis of 3 with 1 M HCl released \( \alpha-L-glucose \) and \( \beta-L-rhamnose \) moieties, while alkaline hydrolysis afforded a ferulic acid moiety, identified by HPLC comparison with authentic samples. The \( ^1H \)-NMR spectrum of 3 exhibited resonances for anomic protons of O-linked sugars at \( \delta_1 \) 4.98 (1H, d, \( J = 7.4 \) Hz) and 5.51 (1H, d, \( J = 7.6 \) Hz), corresponding to two \( \alpha-L-glucoses \), and 4.38 (1H, d, \( J = 1.3 \) Hz) and 5.14 (1H, d, \( J = 1.1 \) Hz), together with typical doublets of two methyl groups at \( \delta_1 \) 0.88 (3H, d, \( J = 6.2 \) Hz) and 0.96 (3H, d, \( J = 6.2 \) Hz), corresponding to two \( \beta-L-rhamnose \) moieties. Full assignment of the \( ^1H \) and \( ^13C \)-NMR data of each sugar was achieved using standard 2D NMR experiments (Tables 1 and 2). The configurations of these sugars were identified from the magnitude of their \( J_1 \) coupling constants and the chemical shifts as \( \beta-L-glucopyranose \) and \( \alpha-L-rhamnopyranose \). The glycosidation sites on the kaempferol aglycone of 3 were determined from the long-range correlations exhibited in the HMBC spectrum from H-1” \( \delta_1 \) 5.51 to C-3 (\( \delta_1 \) 134.6), and from H-1” \( \delta_1 \) 4.98 to C-7 (164.4). Both C-2” and C-6” of the \( \beta-glucose \) attached at C-3 were shifted in the \( ^13C \)-NMR spectrum (Table 2) to \( \delta_1 \) 79.7 and 68.3, respectively, and glycosylated by two \( \alpha-L-rhamnopyranosyl \) moieties (Rha I and Rha II, respectively), according to the
HMBC correlations (Figure 2), their connectivities were detected by the cross peaks between H-2” (δH 3.52) and C-1”” of Rha I (δC 102.5), and between H-2’” (δH 3.27 and 3.71) and C-1”” of Rha II (δC 102.2). Furthermore, the HMBC correlations from significant downfield shifted protons of H-13p were determined from the magnitude of the cross peaks between H-2 and H-6. The results showed that compounds 3 and 4 exhibited as kaempferol 3-O-[2,6-di-O-α-L-rhamnopyranosyl]-β-D-glucopyranoside-7-O-[6-O-(trans-feruloyl)]-β-D-glucopyranoside.

2.1.4. Chemical Structure of Compound 4

Compound 4 was isolated as a yellow amorphous powder, with a negative optical rotation ([α]D 22 81.6). Its molecular formula was established as C48H56O26 by its positive HR-ESI-MS spectrum (m/z 1071.2925 [M + Na]+, calcd for C48H56O26Na 1071.2952), suggesting the lack of one methoxy group compared to 3. Analysis of 2D experiments of 4 including COSY, HSQC and HMBC, and comparison of its 1H- and 13C-NMR values with those of 3 (Tables 1 and 2) showed that 4 was a kaempferol tetracylcoside acylated with a cinnamic acid derivative. The 1H- and 13C-NMR spectra of 4 revealed an identical glycosylation profile to that of 3. Comparison of the HMBC correlations of 4 with those of 3 (Figure 2) detected the same pattern of connectivity defining the linkages between the glycosyl moieties and the kaempferol residue, and between the glycosyl moieties themselves. Thus, the difference between 3 and 4 must be located at the acyl substituent. In particular, the 1H-NMR spectrum (Table 1) showed the typical coupling pattern (AA’BB’ system) of a 1,4-disubstituted aromatic moiety at δH 6.66 and 7.25 (each 2H, d, J = 8.6 Hz) and two trans-coupled double bond protons at δH 6.24 and 7.50 (each 1H, br d, J = 15.9 Hz), as well as to an ester carbonyl (δC 169.1) in the 13C-NMR spectrum assignable to a p-coumaroyl unit. In the HMBC spectrum (Figure 2), the cross peaks between H-2’’” (δH 4.18 and 4.54) and C-9’’’’’ (δC 169.1) placed the p-coumaroyl unit at C-6’’’. The sugar units were identified as D-glucose and L-rhamnose by acid hydrolysis of 4, followed by HPLC analysis with chiral detector in comparison with authentic samples. Anomeric configurations were determined from the magnitude of the J1,2 coupling constants in the 1H-NMR spectrum. Therefore, the structure of 4 was a trans-p-coumaroylated kaempferol tetracylcoside, i.e., kaempferol 3-O-[2,6-di-O-α-L-rhamnopyranosyl]-β-D-glucopyranoside-7-O-[6-O-(trans-p-coumaroyl)]-β-D-glucopyranoside.

2.2. Biological Activities of the Isolated Compounds

Isolated compounds 1-20 from the n-BuOH fraction of C. rotundifolia were evaluated for DPPH radical scavenging assay activity (Table 3) using Trolox as a positive control (IC50: 29.2 ± 0.39 µM). The results showed that compounds 17 and 18 exhibited potent radical scavenging activity with IC50 values of 14.5 and 11.7 µM, respectively. On the other hand, compounds 15 and 16 showed moderate activity with IC50 values of 43.0 and 31.8 µM, respectively. These results are consistent with previous studies on the effectiveness of myricetin as a potent antioxidant agent which could prevent and slow the progress of ageing or various diseases associated with oxidative stress.

<table>
<thead>
<tr>
<th>Isolated Compounds</th>
<th>DPPH (IC50, µM)</th>
<th>AGEs (IC50, µM)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Kaempferol 3-O-[2,6-di-O-α-L-rhamnopyranosyl]-β-D-glucoside (7)</td>
<td>&gt;100</td>
<td>85.5 ± 3.5</td>
</tr>
<tr>
<td>Myricetin 3-O-[2,6-di-O-α-L-rhamnopyranosyl]-β-D-glucoside (15)</td>
<td>43.0 ± 1.15</td>
<td>&gt;100</td>
</tr>
<tr>
<td>Myricetin 3-O-β-neohesperidoside (16)</td>
<td>31.8 ± 0.63</td>
<td>&gt;100</td>
</tr>
<tr>
<td>Myricetin 3-O-β-D-glucoside (17)</td>
<td>14.5 ± 1.15</td>
<td>96.5 ± 1.8</td>
</tr>
<tr>
<td>Myricetin (18)</td>
<td>11.7 ± 1.8</td>
<td>34.9 ± 1.2</td>
</tr>
<tr>
<td>Aminoguanidine hydrochloride</td>
<td>29.2 ± 0.39</td>
<td>n.d.</td>
</tr>
<tr>
<td>Trolox</td>
<td>7818 ± 34.4</td>
<td>n.d.</td>
</tr>
</tbody>
</table>

n.d.: not determined.
Inhibitory effects of isolated compounds (1–20) on AGEs formation were also evaluated using a fluorescence method [14]. As shown in Table 3, compound 7 (IC\textsubscript{50}: 85.5 ± 3.5 µM), 17 (IC\textsubscript{50}: 96.5 ± 1.8 µM) and 18 (IC\textsubscript{50}: 34.9 ± 1.2 µM) showed strong inhibitory activity than that of a positive control, aminoguanidine hydrochloride (IC\textsubscript{50}: 7818 ± 34.4 µM). The results revealed that the most inhibitory effect was exhibited by myricetin (18). Comparison of IC\textsubscript{50} values of myricetin (18) and its tri-, di-, and mono-glycosides (15, 16, and 17, respectively) indicated that the number of sugar moieties at C-3 of myricetin aglycone decreases the inhibition activity of AGEs formation. The other compounds did not show any inhibitory activity.

3. Materials and Methods

3.1. General Methods

Optical rotations were measured on a JASCO P-1030 polarimeter (Jasco, Tokyo, Japan). UV and IR spectra were obtained on a Jasco V-520 UV/Vis spectrophotometer and a Horiba FT-710 Fourier transform infrared spectrophotometer (Horiba, Kyoto, Japan), respectively. NMR experiments were measured on a Bruker Avance 500, 600 and 700 MHz spectrometers (Bruker, Billerica, MA, USA), with tetramethylsilane (TMS) as an internal standard. Positive ion HR-ESI-MS spectra were recorded using a LTQ Orbitrap XL mass spectrometer (Thermo Fisher Scientific, Waltham, MA, USA). Column chromatography (CC) was performed on Diaion HP-20 (Mitsubishi Chemical Corp., Tokyo, Japan), silica gel 60 (230–400 mesh, Merck, Darmstadt, Germany), and octadecyl silica (ODS) gel (Cosmosil 75C\textsubscript{18}-OPN (Nacalai Tesque, Kyoto, Japan; \( \Phi = 35 \) mm, \( L = 350 \) mm), and TLC was performed on precoated silica gel plates 60 GF\textsubscript{254} (0.25 mm in thickness, Merck). HPLC was performed on ODS gel (Inertsil ODS-3, GL-science, 10 mm × 250 mm, flow rate 2.5 mL/min) with a mixture of H\textsubscript{2}O, acetone and MeOH, and the eluate was monitored by refractive index and/or a UV detector. An amino column (Shodex Asahipak NH2P-50 4E (4.6 mm × 250 mm), CH\textsubscript{3}CN-H\textsubscript{2}O (3:1) 1 mL/min) together with a chiral detector (Jasco OR-2090plus) was used for HPLC analysis of sugars obtained after hydrolysis [15].

3.2. Plant Material

Aerial parts of \textit{Cadaba rotundifolia} were collected from eastern region, near Makkah, Saudi Arabia, in April 2017. The collected plant was identified and classified by a plant taxonomist at Biology Department–Umm Al-Qura University, Makkah, Saudi Arabia. A voucher specimen (No. 15501) was deposited in the Herbarium of the Department of Pharmacognosy, Pharmacy College, King Saud University.

3.3. Extraction and Isolation

The air-dried powdered aerial parts (6.0 Kg) of \textit{C. rotundifolia} Forssk. were extracted with MeOH (10 L × 7) and then concentrated under reduced pressure to give a viscous gummy material (800 g). This residue was suspended in 900 mL of distilled water and fractionated by shaking with petroleum ether (bp. 60–80 °C) (3L × 5, 132.9 g), CH\textsubscript{2}Cl\textsubscript{2} (3 L × 5, 6 g), EtOAc (3 L × 5, 10.9 g) and n-BuOH (3 L × 5, 118 g), respectively. Part of \textit{n}-BuOH fraction (112 g) was fractionated by Diaion HP-20 column chromatography (\( \Phi = 35 \) mm, \( L = 350 \) mm, 900 g). The column was eluted initially with H\textsubscript{2}O, then with a MeOH/H\textsubscript{2}O stepwise gradient with increasing MeOH content (20, 40, 60, 80, 100% MeOH, respectively, 3 L each), yielding five fractions (Frs. Cr1–Cr5). The fraction Cr2 (10.7 g) was subjected to silica gel column chromatography (\( \Phi = 6 \) cm, \( L = 35 \) cm, 900 g). The column was eluted initially with H\textsubscript{2}O, then with a MeOH/H\textsubscript{2}O stepwise gradient with increasing MeOH content (20, 40, 60, 80, 100% MeOH, respectively, 3 L each), yielding five fractions (Frs. Cr1–Cr5). The fraction Cr2 (10.7 g) was subjected to silica gel column chromatography (\( \Phi = 3 \) cm, \( L = 15 \) cm, 100 g) eluting with increasing amount of MeOH in CHCl\textsubscript{3} (20:1, 10:1, 7:1, 5:1, 3:1, 1:1, finally with 100% MeOH, 500 mL each) yielding seven fractions (Frs. Cr2.1–Cr2.7). Fraction Cr2.4 (3.5 g) was subjected to open reversed phase (ODS) column chromatography with a linear gradient of 10% acq. MeOH (400 mL–100% methanol (400 mL), affording eight fractions (Frs. Cr2.4.1–Cr2.4.8). The residue of fraction Cr2.4.4 (187 mg) was purified by preparative HPLC using 46% acq. acetone to give 18 (5.0 mg) and 20 (25.1 mg). Fraction Cr2.5 (2.4 g) was subjected to ODS column chromatography eluting with a linear gradient of 10% acq. MeOH.
(400 mL)–100% MeOH (400 mL), to give seven fractions (Frs. Cr2.5.1–Cr2.5.7). The residue of fraction Cr2.5.4 (554 mg) was purified by preparative HPLC (25% aq. acetone) to give 6 (4.2 mg). Fraction Cr2.7 (2.6 g) was subjected to ODS column chromatography (linear gradient of 10% aq. MeOH (400 mL)–100% MeOH (400 mL), giving six fractions (Frs. Cr2.7.1–Cr2.7.6). The residue of fraction Cr2.7.2 (632 mg) was purified by preparative HPLC (25% aq. acetone) to give 15 (5.6 mg) and 16 (11.0 mg). Then residue of fraction Cr2.7.3 (1.2 g) was also purified by preparative HPLC (28% aq. acetone) to give 7 (17.3 mg) and 19 (9.2 mg). Fraction Cr3 (15.5 g) was subjected to silica gel column chromatography (Φ = 3 cm, L = 20 cm, 150 g) with increasing amounts of MeOH in CHCl3 (20:1, 10:1, 7:1, 5:1, 3:1, 1:1, then 100% MeOH, 400 mL each), yielding seven fractions (Frs. Cr3.1–Cr3.7). Fraction Cr3.3 (2.9 g) was subjected to ODS column chromatography (linear gradient of 10% aq. MeOH (400 mL)–100% MeOH (400 mL)), affording nine fractions (Frs. Cr3.3.1–Cr3.3.9). The residue of fraction Cr3.3.4 (645 mg) was purified by preparative HPLC (28% aq. acetone) to give 10 (30.1 mg). Fraction Cr3.4 (2.8 g) was subjected to ODS column chromatography (linear gradient of 10% aq. MeOH (400 mL)–100% MeOH (400 mL)) leading to seven fractions (Frs. Cr3.4.1–Cr3.4.7). The residue of fraction Cr3.4.2 (1.2 g) was purified by preparative HPLC (28% aq. acetone) to give 17 (15.2 mg). The residue of fraction Cr3.4.4 (187 mg) was also purified by preparative HPLC (40% aq. acetone) to give 1 (7.4 mg) and 5 (2.0 mg). Fraction Cr3.5 (4.4 g) was subjected to ODS column chromatography (linear gradient of 10% aq. MeOH (400 mL)–100% MeOH (400 mL)) providing seven fractions (Frs. Cr3.5.1–Cr3.5.7). The residue of fraction Cr3.5.4 (692 mg) was purified by preparative HPLC (28% aq. acetone) to give 8 (39.0 mg) and 9 (20.0 mg). The other residue of fraction Cr3.5.5 (65.7 mg) was purified by preparative HPLC (32% aq. acetone) to give 11 (3.0 mg). Fraction Cr3.6 (760 mg) was subjected to ODS column chromatography (linear gradient of 10% aq. MeOH (400 mL)–100% MeOH (400 mL)), affording seven fractions (Frs. Cr3.6.1–Cr3.6.7). The residue of fraction Cr3.6.4 (923 mg) was purified by preparative HPLC (25% aq. acetone). The following compounds were separated: 2 (5.0 mg), 12 (12.0 mg), 13 (2.0 mg) and 14 (10.0 mg). The residue of fraction Cr3.6.5 (963 mg) was purified by preparative HPLC (35% aq. acetone) to give 3 (7.8 mg) and 4 (2.0 mg).

The known compounds 5–20 were identified by comparison of their spectroscopic data with those reported in the literature as kaempferol 3-O-[2-O-(trans-p-coumaroyl)-3-O-β-d-glucopyranosyl]-β-d-glucopyranoside (5) [12], kaempferol 3,4′-di-O-β-d-glucoside (6) [16], kaempferol 3-O-[2,6-di-O-α-L-rhamnopyranosyl]-β-d-glucoside (7) [17], kaempferol 3-O-β-neohesperidoside (8) [17], kaempferol 3-O-β-rutinoside (9) [17], kaempferol 3-O-β-d-glucoside (10) [17], rhamnocitrin 3-O-β-neohesperidoside (11) [18], kaempferol 3-O-β-neohesperidoside-7-O-[2-O-(trans-p-coumaroyl)]-3-O-β-d-glucopyranosyl]-β-d-glucopyranoside (12) [13], kaempferol 3-O-β-neohesperidoside-7-O-[2-O-(trans-feruloyl)]-β-d-glucopyranoside (13) [13], kaempferol 3-O-β-neohesperidoside-7-O-[2-O-(trans-p-coumaroyl)]-β-d-glucopyranoside (14) [13], myricetin 3-O-[2,6-di-O-α-L-rhamnopyranosyl]-β-d-glucoside (15) [17], myricetin 3-O-β-neohesperidoside (16) [17], myricetin 3-O-β-D-glucoside (17) [17], myricetin (18) [19], beitingxuahuangtong C (19) [20], phloretin (20) [21].

3.4. Spectroscopic Data of Compounds 1–4

Kaempferol 3-O-[2-O-(trans-feruloyl)]-3-O-β-d-glucopyranosyl]-β-d-glucopyranoside (1)

Yellow amorphous powder [α]D 22 +79.0 (c 0.42, MeOH); HR-ESI-MS: m/z: 809.1888 [M + Na]+ (calcd for C37H38O19Na 809.1899); UV λmax (MeOH) nm (log ε): 327 (4.21), 299 (4.12), 267 (4.13), 231 (4.23); IR (film) νmax: 3388, 2936, 1710, 1658, 1604, 1513, 1446, 1363, 1176, 1070, 1028 cm−1; 1H and 13C data see Tables 1 and 2.

Kaempferol 3-O-β-neohesperidoside-7-O-[2-O-(cis-p-coumaroyl)]-3-O-β-d-glucopyranosyl]-β-d-glucopyranoside (2)

Yellow amorphous powder [α]D 22 +14.5 (c 0.60, MeOH); HR-ESI-MS: m/z: 1087.2875 [M + Na]+ (calcd for C48H56O27Na 1087.2901); UV λmax (MeOH) nm (log ε): 351 (3.97), 316 (4.14), 267 (4.04), 229...
(3.99); IR (film) ν_{max}: 3397, 2924, 1721, 1657, 1512, 1442, 1367, 1157, 1072, 1024 cm⁻¹; \(^1^H\) and \(^1^3^C\) data see Tables 1 and 2.

**Kaempferol 3-O-[2,6-di-O-\(\alpha\)-l-rhamnopyranosyl]-\(\beta\)-d-glucopyranoside-7-O-[6-O-(trans-feruloyl)]-\(\beta\)-d-glucopyranoside (3)**

Yellow amorphous powder [\(\alpha\)]\textsubscript{22}D = −82.5 (c 1.02, MeOH); HR-ESI-MS: \(m/z\): 1101.3026 [M + Na]\(^+\) (calcd for C\(_{49}\)H\(_{58}\)O\(_{27}\)Na 1101.3058); UV λ\(_{max}\) (MeOH) nm (log \(ε\)): 329 (4.17), 295 (4.00), 267 (4.05), 233 (4.11); IR (film) ν\(_{max}\): 3389, 2932, 1712, 1651, 1601, 1513, 1455, 1371, 1180, 1071, 1024 cm⁻¹; \(^1^H\) and \(^1^3^C\) data see Tables 1 and 2.

**Kaempferol 3-O-[2,6-di-O-\(\alpha\)-l-rhamnopyranosyl]-\(\beta\)-d-glucopyranoside-7-O-[6-O-(trans-p-coumaroyl)]-\(\beta\)-d-glucopyranoside (4)**

Yellow amorphous powder [\(\alpha\)]\textsubscript{22}D = −81.6 (c 0.36, MeOH); HR-ESI-MS: \(m/z\): 1071.2925 [M + Na]\(^+\) (calcd for C\(_{48}\)H\(_{56}\)O\(_{26}\)Na 1071.2952); UV λ\(_{max}\) (MeOH) nm (log \(ε\)): 348 (4.01), 317 (4.15), 267 (4.05), 228 (4.09); IR (film) ν\(_{max}\): 3379, 2912, 1660, 1499, 1445, 1372, 1172, 1074, 1034 cm⁻¹; \(^1^H\) and \(^1^3^C\) data see Tables 1 and 2.

### 3.5. Acid Hydrolysis

The isolated compounds 1-4 (1.0 mg each) were hydrolyzed with 1 M HCl (1.0 mL) at 80 °C for 3 h, then cooled. The reaction mixtures were neutralized with Amberlite IRA96SB (OH⁻ form), then filtered to remove the resin. The filtrates were extracted with EtOAc. The aqueous layers were analyzed by HPLC with an amino column [Asahipak NH2P-50 4E, CH\(_3\)CN-H\(_2\)O (3:1), 1 mL/min] and a chiral detector (JASCO OR-2090plus) in comparison with authentic standards of D-glucose and L-rhamnose. The aqueous layers of the hydrolyzed compounds showed a peak at \(t_R\) 8.15 min which coincided with that of D-glucose (positive optical rotation). The \(t_R\) of the L-rhamnose (negative optical rotation) was 6.5 min [15].

### 3.6. Alkaline Hydrolysis

Solutions of 1-3 (3.0 mg each) in 50% aqueous 1,4-dioxane (0.5 mL) were treated with 10% aqueous KOH (0.5 mL) and stirred at 37 °C for 1 h. The reaction mixtures were neutralized with an ion-exchange resin Amberlite IR-120B (H\(^+\)-form), then filtered and evaporated. The residues were dissolved in EtOAc and subjected to HPLC analysis [Cosmosil C18-PAQ, MeOH-H\(_2\)O (45:50), 1 mL/min] with UV (245 nm) detection to identify ferulic acid (\(t_R\) 16.7 min) from 1 and 3, and \(p\)-coumaric acid (\(t_R\) 15.1 min) from 2 [22]. The authentic cis-\(p\)-coumaric acid was prepared according to the literature [23].

### 3.7. DPPH Radical Scavenging Activity

Free radical scavenging activity were evaluated by using a quantitative DPPH assay. The absorbance of various concentrations of test compounds dissolved in 100 \(\mu\)L of MeOH in 96-well microtiter plate were measured at 515 nm as (A\(_{\text{blank}}\)). Then, a 100 \(\mu\)L of DPPH solution (200 \(\mu\)M) was added to each well. The plate was incubated in dark chamber at room temperature for 30 min before measuring the absorbance (A\(_{\text{sample}}\)) again. The following equation was used to calculate % inhibition of free radicals:

\[
\% \text{ Inhibition} = \left[1 - \frac{(A_{\text{sample}} - A_{\text{blank}})}{(A_{\text{control}} - A_{\text{blank}})}\right] \times 100
\]

where A\(_{\text{control}}\) is the absorbance of control (DMSO and all reagents, except the test compound). Their IC\(_{50}\) values were determined based on three independent experiments. Trolox was used as reference compound [24].
3.8. Determination of In Vitro AGEs Formation

The reaction mixture including 0.5 M ribose and 10 mg/mL bovine serum albumin in 50 mM phosphate buffer (pH 7.4)- 0.02% sodium azide was mixed with the test compounds, and incubated for 24 h at 37 °C. The formation of fluorescent AGEs were measured with a fluorometric microplate reader (Ex: 370 nm, Em: 440 nm, EnSpire, PerkinElmer, Inc., Waltham, MA, USA). Experiments were performed in triplicate and IC₅₀ values were calculated by linear regression [14].

4. Conclusions

In conclusion, four new flavonoid glycosides 1–4 together with 16 known compounds 5–20 were isolated from the aerial parts of Cadaba rotundifolia (Forssk.) (Qadab) collected in Saudi Arabia. The chemical structures of the new compounds were elucidated by intensive spectroscopic analyses and chemical reactions. Among the isolated compounds, 15–18 exhibited significant radical scavenging activity and compounds 7, 17 and 18 showed strong inhibitory activity towardageformation. These results suggest that C. rotundifolia and their chemical constituents may ameliorate oxidative stress in various degenerative disorders including Alzheimer’s disease, diabetes, atherosclerosis and aging. Finally, it is noteworthy that most of the known compounds, 5–8, 11–16, 19, and 20, were isolated from the family Capparaceae for the first time. The remaining three known compounds, 9, 10, 17, also have not been isolated from the title plant so far. These results indicated that the characteristic circumstances of the oases surrounded by desert in Saudi Arabia provide promising and interesting natural resources for future drug development.


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Sample Availability: Samples of the compounds 1–4 are available from the authors.