



Original Article

HPLC-MS profiling of the multidrug-resistance modifying resin glycoside content of *Ipomoea alba* seeds



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ABSTRACT

High performance liquid chromatography profiling with mass spectrometry detection was applicable to identify known and novel multidrug-resistance glycolipid inhibitors from the complex resin glycosides mixture of *Ipomoea alba* L., Convolvulaceae, seeds. Albinosides X and XI were purified by recycling liquid chromatography and their structural elucidation was accomplished by nuclear magnetic resonance. Albinoside XI exerted a strong potentiation of vinblastine susceptibility in multidrug-resistant human breast carcinoma cells.

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Introduction

Ipomoea alba L., Convolvulaceae, or moon vine, is a night-blooming morning glory native to tropical Americas (McDonald, 1994). In Mexico, the decoction of the leaves has been used for the treatment of paralysis and swelling of tissues (Lim, 2014). This species is an ornamental plant but it has also been reported its invasive potential (Foxcroft et al., 2008).

Previously, eleven resin glycosides were isolated from moon vine, calonyctins A₁ and A₂ from EtOH-soluble extracts of dried leaves (Fang et al., 1993) and albinosides I-IX from CHCl₃-soluble extracts of seeds (Cruz-Morales et al., 2012, 2016). Resin glycosides are complex mixtures of oligosaccharides of monohydroxy and dihydroxy C-14 and C-16 fatty acids, which represent distinctive secondary metabolites restricted to the morning glory family (Pereda-Miranda et al., 2010). These mixtures of glycolipids are a source of efflux pumps modulators responsible for the multidrug resistant (MDR) phenotype in Gram-positive (Pereda-Miranda et al., 2006) and -negative bacteria (Corona-Castañeda and Pereda-Miranda, 2012), as well as in mammalian cancer cells (Figueroa-González et al., 2012). MDR phenotype is considered a major cause for the failure of anticancer agents. Thus, the use of efflux pump modulators co-administered with cytotoxic drugs results in a susceptibility equivalent to that of a sensitive cell

(Figueroa-González et al., 2012). Hence, in this context, the present HPLC-MS profiling of the resin glycoside content of *Ipomoea alba* seeds was undertaken to identify novel multidrug-resistance glycolipid inhibitors.

Materials and methods

General experimental procedures

The experimental procedures, including HPLC, GC-MS, MS and NMR instrumentation, were previously described (Cruz-Morales et al., 2012, 2016).

Chemicals and cell lines

RPMI 1640 medium and fetal bovine serum were purchased from Gibco (Life Technologies, Carlsbad, CA) and sulforhodamine B (SRB), reserpine, and vinblastine from Sigma-Aldrich (St. Louis, MO). Breast (MCF-7 and MDA-MB-231), cervix (HeLa), and colon (HCT-15 and HCT-116) carcinoma cell lines were acquired from the American Type Culture Collection. The resistant counterpart MCF-7/Vin has been subcultured during seven years (Figueroa-González et al., 2012).

Plant material

Moon vine seeds (*Ipomoea alba* L., Convolvulaceae; item # 01052-PK-P1) were purchased from Park Seed (Greenwood, SC)

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in March 2015. Eight seeds were germinated and two seedlings were grown to maturity under the conditions previously described (Cruz-Morales et al., 2016). Vouchers were deposited at the Arkansas State University Herbarium (STAR 027009).

Extraction and isolation

Maceration of dried powdered seeds (300 g) was performed at room temperature with CHCl_3 . Precipitation of the resin glycoside with MeOH was achieved from the CHCl_3 -soluble extract to obtain a white solid (4.5 g), which was washed with hexane. The precipitated solid was analyzed by HPLC using a reversed-phase Symmetry C_{18} column (Waters; 7 μm , 19 \times 300 mm), an isocratic elution with MeOH– CH_3CN – H_2O (5:4:1), and a flow rate of 4 ml/min with a sample injection of 500 μl (concentration: 0.1 mg/ μl). The resulting chromatogram showed eleven resolved peaks (Fig. 1). Eluates with t_R values of 83.3 min (20.9 mg) and 147.1 min (18.3 mg) were collected by the heart cutting technique (Pereda-Miranda and Hernández-Carlos, 2002). Each subfraction was independently reinjected and purified by recycling HPLC to achieve homogeneity after ten consecutive cycles by application of peak shaving technique (Pereda-Miranda and Hernández-Carlos, 2002) until overlapped minor impurities were separated: Symmetry C_{18} column (Waters; 7 μm , 19 \times 300 mm) with an elution of MeOH– CH_3CN – H_2O (10:7:3) and a flow rate of 6 ml/min. These procedures allowed the purification of **1** (15.5 mg; t_R 8.6 min) and **2** (13.5 mg; t_R 10.7 min).

Albinoside X (1): White solid; mp 128–131 °C; $[\alpha]_{589}^{20}$ –11.7, $[\alpha]_{578}^{20}$ –12.0, $[\alpha]_{546}^{20}$ –13.7, $[\alpha]_{436}^{20}$ –20.6, $[\alpha]_{365}^{20}$ –26.2 (c 1.0, MeOH); ^1H and ^{13}C NMR see Table 2 positive ESIMS: m/z 997 $[\text{M}+\text{Na}]^+$; negative FABMS: m/z 973 $[\text{M}-\text{H}]^-$, 891 $[\text{M}-\text{H}-\text{C}_5\text{H}_6\text{O}$ (tigloyl)] $^-$, 809 $[\text{M}-\text{H}-\text{C}_5\text{H}_6\text{O}$ (tigloyl)] $^-$, 663 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 517 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 389 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 243 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$; HRESIMS m/z 973.5081 $[\text{M}-\text{H}]^-$ (calcd for $\text{C}_{48}\text{H}_{77}\text{O}_{20}$ requires 973.5014).

Albinoside XI (2): White solid; mp 138–140 °C; $[\alpha]_{589}^{20}$ –13.3, $[\alpha]_{578}^{20}$ –13.8, $[\alpha]_{546}^{20}$ –15.7, $[\alpha]_{436}^{20}$ –22.9, $[\alpha]_{365}^{20}$ –29.5 (c 1.0, MeOH); ^1H and ^{13}C NMR see Table 2; positive ESIMS: m/z 1025 $[\text{M}+\text{Na}]^+$; negative FABMS: m/z 1001 $[\text{M}-\text{H}]^-$, 919 $[\text{M}-\text{H}-\text{C}_5\text{H}_6\text{O}$ (tigloyl)] $^-$, 837 $[\text{M}-\text{H}-\text{C}_5\text{H}_6\text{O}$ (tigloyl)] $^-$, 691 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 545 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 417 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 271 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$; HRESIMS m/z 1001.5392 $[\text{M}-\text{H}]^-$ (calcd for $\text{C}_{50}\text{H}_{81}\text{O}_{20}$ requires 1001.5327).

Alkaline hydrolysis of compounds 1–2

Solutions of compounds **1** and **2** (10 mg for each sample), in 5% KOH– H_2O (1 ml) were refluxed at 95 °C for 3 h. Then, the mixtures were acidified to pH 5.0 and extracted with CHCl_3 (2 \times 2.5 ml) and Et_2O (2 \times 2.5 ml). The CHCl_3 -soluble layer was washed, dried over anhydrous Na_2SO_4 , evaporated, and analyzed by GC–MS. Tiglic acid was detected as the only volatile product for compounds **1** and **2** (t_R 6.95 min); m/z $[\text{M}]^+$ 100 (30), 73 (100), 55 (22). The aqueous layer was extracted with *n*-BuOH (10 ml) and concentrated. Hydrolysis of compound **1** afforded **12** (4.5 mg), and compound **2** yielded **13** (4.3 mg).

Albinosinic acid H (12): white solid; mp 96–98 °C; $[\alpha]_{\text{D}}^{20}$ –32.1 (c 1.0, MeOH); negative ESI-MS m/z 827 $[\text{M}-\text{H}]^-$. FABMS m/z 827 $[\text{M}-\text{H}]^-$, 681 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 535 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 389 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, and 243 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit), aglycone] $^-$.

Albinosinic acid I (13): white solid; mp 102–105 °C; $[\alpha]_{\text{D}}^{20}$ –30.6 (c 1.0, MeOH); negative ESI-MS m/z 855 $[\text{M}-\text{H}]^-$. FABMS m/z

855 $[\text{M}-\text{H}]^-$, 709 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 563 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, 417 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit)] $^-$, and 271 $[\text{M}-\text{H}-\text{C}_6\text{H}_{10}\text{O}_4$ (methylpentose unit), aglycone] $^-$.

Sugar analysis

Compounds **12** and **13** (3 mg of each one) in 10 ml HCl 4N were independently refluxed at 90 °C for 1 h. Then, mixtures were diluted with 2.5 mL H_2O and extracted with ether (2 \times 5 ml). The organic layer was evaporated, re-suspended in CHCl_3 (3 ml), and alkylated with CH_2N_2 . The aqueous layer was neutralized with KOH 1N and extracted with *n*-BuOH (10 ml), then washed with H_2O (2 \times 5 ml) and concentrated. The thiazolidine derivatives of each mixture were prepared according to previously described procedures (Cruz-Morales et al., 2016). These mixtures were treated with chlorotrimethylsilane (Sigma Sil-A) and analyzed by GC-MS (Cruz-Morales et al., 2016).

Identification of aglycones

The derivatized (CH_2N_2) residues obtained from acid hydrolysis of **12** and **13** were analyzed by normal-phase HPLC ($\mu\text{Porasil}$, 3.9 \times 300 mm, 10 μm), using an elution [hexane– CHCl_3 – Me_2CO (6:3:1)] and a flow rate of 1 ml/min to identify methyl (11S)-hydroxytetradecanoate (t_R 18.0 min) from compound **1** and methyl (11S)-hydroxyhexadecanoate (t_R 15.5 min) from compound **2** by comparison with authentic samples (Cruz-Morales et al., 2016).

Cytotoxicity and modulation of multidrug-resistance assays

Cytotoxicity of compounds **1** and **2** was determined by using the SRB assay. Cell lines were maintained in RPMI 1640 medium supplemented with 10% fetal bovine serum and cultured at 37 °C in 5% CO_2 in air (100% humidity). MCF-7/Vin $^+$ cells were cultured in medium containing 0.192 $\mu\text{g}/\text{ml}$ vinblastine. A stock of MCF-7/Vin cells was also maintained in vinblastine-free medium (MCF-7/Vin $^-$). Cells at log phase were treated in triplicate with test samples (0.2–25 $\mu\text{g}/\text{ml}$) and incubated for 72 h. For the reversal effects, sensitive MCF-7 and MDR MCF-7/Vin cells were seeded into 96-well plates and treated with various concentrations of vinblastine (0.00064–10 $\mu\text{g}/\text{ml}$) in the presence or absence of glycolipids (25 $\mu\text{g}/\text{ml}$) for 72 h. The ability of glycolipids to potentiate vinblastine cytotoxicity was measured by calculating the IC_{50} (Figueroa-González et al., 2012).

Results and discussion

HPLC profiling of the resin glycoside content and isolation of 1 and 2

Preparative reversed-phase HPLC-RI (refractive index detection), using the heart-cutting technique (Pereda-Miranda and Hernández-Carlos, 2002), was used to separate eleven major constituents from the resin glycoside fraction of moon vine seeds (Fig. 1). Profiling of this resin glycosides based on MS allowed dereplication to be performed after fractionation, which included drying and redissolution of the eluted fractions (Potterat and Hamburger, 2013) in order to identify known and novel compounds (Smyth et al., 2012). Each collected peak was analyzed by ESIMS and FAB-MS in both positive and negative modes. Albinosides I–X (3–11) were unambiguously identified by comparison of the fragmentation ions observed in negative mode FAB-MS (Cruz-Morales et al., 2012, 2016), as well as the adduct ion $[\text{M}+\text{Na}]^+$ detected in ESI (Table 1). This HPLC profiling indicated the presence of

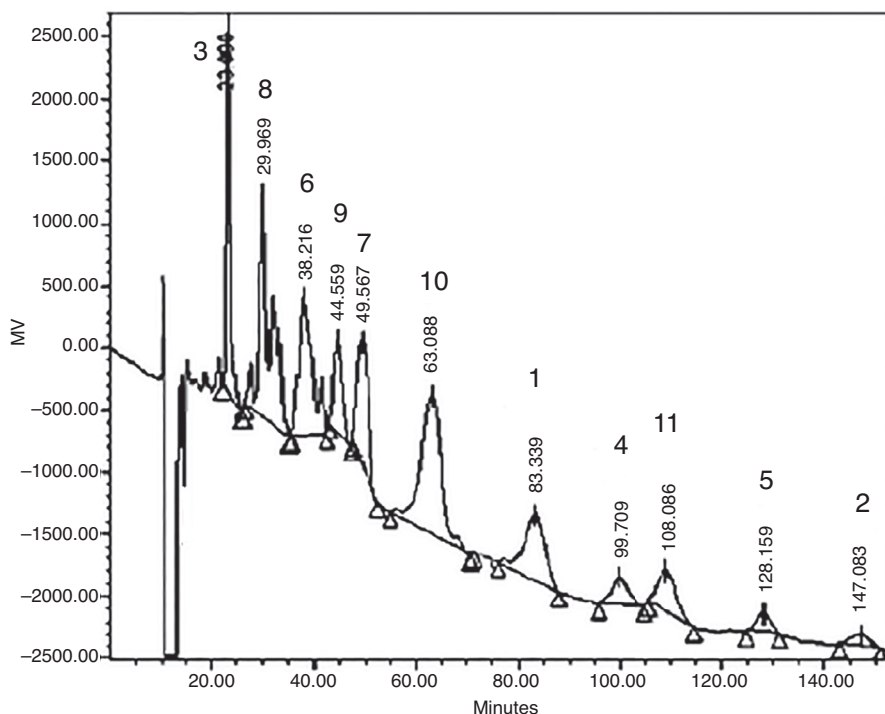


Fig. 1. HPLC chromatogram for the resin glycoside mixture from CHCl_3 -soluble extract. Instrumental conditions: Symmetry C_{18} column (waters: 19×300 mm, 7 mm); isocratic elution with $\text{MeOH}-\text{CH}_3\text{CN}-\text{H}_2\text{O}$ (10:7:3); flow rate: 4.0 ml/min; sample injection: 500 μl [0.1 mg/ μl]. Compounds: **1** ($t_R = 83.3$ min, albinoside X); **2** ($t_R = 147.1$ min, albinoside XI); **3** ($t_R = 23.1$ min, albinoside I); **4** ($t_R = 99.7$ min, albinoside II); **5** ($t_R = 128.2$ min, albinoside III); **6** ($t_R = 38.2$ min, albinoside IV); **7** ($t_R = 49.6$ min, albinoside V); **8** ($t_R = 29.9$ min, albinoside VI); **9** ($t_R = 44.6$ min, albinoside VII); **10** ($t_R = 63.1$ min, albinoside VIII); **11** ($t_R = 109.1$ min, albinoside IX).

two previously unknown glycolipids with $t_R = 83.3$ and 147.1 min (Fig. 1), which were named as albinoside X (**1**) and albinoside XI (**2**), respectively. Each new peak was recycled in HPLC to achieve chromatographic homogeneity, allowing the purification of compounds **1** and **2**.

Structural elucidation of **1** and **2**

Albinoside X (**1**) presented a deprotonated molecule ion peak at m/z 973.5081 ($[\text{M}-\text{H}]^-$; $\text{C}_{48}\text{H}_{77}\text{O}_{20}$, calcd error: $\delta = -0.80$ ppm) in HRESIMS, while albinoside XI (**2**) showed a $[\text{M}-\text{H}]^-$ peak at m/z 1001.5392 ($\text{C}_{50}\text{H}_{81}\text{O}_{20}$, calcd error: $\delta = -0.70$ ppm). The difference of 28 mass units between **1** and **2**, and the observation of an identical pattern for the glycosidic cleavages, allowed to recognize the same linear tetrasaccharide core in both natural products. The presence of 11S-hydroxytetradecanoic acid as the aglycone for compound **1** was confirmed by the diagnostic FAB-MS negative ions at m/z 827 $[\text{M}-\text{H}]^-$, 681, 535, 389 and 243 (aglycone), which corresponded to four consecutive eliminations of methyl pentose units ($\text{C}_6\text{H}_{10}\text{O}_4$, 146 uma), for its saponification derivative, albinosinic acid H (**12**); while for the albinosinic acid I (**13**), negative ions at

m/z 855 $[\text{M}-\text{H}]^-$, 709, 563, 417, and 271 confirmed the aglycone as 11S-hydroxyhexadecanoic acid for natural product **2**.

Saponification of both natural products liberated 2-methyl-2-butenic acid as the acylating group. The same sugar composition was obtained by acid hydrolysis of both compounds **12** and **13**, L-rhamnose, D-quinovose, and D-fucose in the ratio 2:1:1. The NMR spectra of both natural products **1** and **2** were very similar (Table 2). For **1**, four anomeric protons were observed at δ_H 4.77 (1H, d, $J = 7.6$ Hz; δ_C 103.2, Qui-1); 5.89 (1H, brs; δ_C 100.4, Rha-1); 5.29 (1H, d, $J = 7.6$ Hz; δ_C 107.1, Fuc-1), 5.64 (1H, brs; δ_C 99.9, Rha'-1) in the HSQC spectra. The HSQC spectra registered for compound **2** was almost identical with four anomeric signal centered at δ_H 4.75 (1H, d, $J = 7.6$ Hz; δ_C 103.0, Qui-1); 5.88 (1H, brs; δ_C 100.2, Rha-1); 5.30 (1H, d, $J = 7.6$ Hz; δ_C 106.9, Fuc-1), 5.64 (1H, brs; δ_C 99.8, Rha'-1). The glycosylation sequence was established by the long-range heteronuclear coupling correlations ($^3J_{\text{CH}}$) in the HMBC spectra. For instance, key correlations were observed in compounds **1** and **2**: (a) the connectivity between H-1 of the quinovose (**1**: δ_H 4.77; **2**: δ_H 4.75) and C-11 of the fatty acid (**1**: δ_C 81.7; **2**: δ_C 80.9); (b) H-2 of quinovose (**1**: δ_H 4.44; **2**: δ_H 4.45) with C-1 of rhamnose (**1**: δ_C 100.4; **2**: δ_C 100.2); (c) H-4 of rhamnose (**1**:

Table 1
Albinosides identification with ESI-MS and FAB-MS fragmentation.

Compound	t_R (min)	Formula	MW	ESI $[\text{M} + \text{Na}]^+$	FAB $[\text{M}-\text{H}]^-$	Fragment ions (FAB negative mode, m/z)
1 (Albinoside X)	83.3	$\text{C}_{48}\text{H}_{78}\text{O}_{20}$	974.508	997	973	891, 809, 663, 517, 389, 243
2 (Albinoside XI)	147.1	$\text{C}_{50}\text{H}_{82}\text{O}_{20}$	1002.539	1025	1001	919, 837, 691, 545, 417, 271
3 (Albinoside I)	23.1	$\text{C}_{51}\text{H}_{86}\text{O}_{26}$	1114.532	1137	1113	1013, 971, 825, 679, 533, 389, 243
4 (Albinoside II)	99.7	$\text{C}_{54}\text{H}_{88}\text{O}_{24}$	1120.558	1153	1119	1037, 891, 745, 517, 389, 243
5 (Albinoside III)	128.2	$\text{C}_{57}\text{H}_{96}\text{O}_{24}$	1164.620	1187	1163	1081, 935, 853, 707, 561, 417, 271
6 (Albinoside IV)	38.2	$\text{C}_{49}\text{H}_{82}\text{O}_{24}$	1054.513	1077	1053	971, 907, 825, 679, 533, 389, 243
7 (Albinoside V)	49.6	$\text{C}_{54}\text{H}_{90}\text{O}_{25}$	1138.569	1161	1137	1037, 891, 809, 791, 745, 663, 517, 389, 243
8 (Albinoside VI)	29.9	$\text{C}_{45}\text{H}_{76}\text{O}_{22}$	968.470	991	967	867, 679, 551, 533, 389, 243
9 (Albinoside VII)	44.6	$\text{C}_{45}\text{H}_{76}\text{O}_{21}$	952.477	975	951	851, 809, 663, 535, 517, 389, 371, 243
10 (Albinoside VIII)	63.1	$\text{C}_{48}\text{H}_{78}\text{O}_{21}$	990.496	1013	989	907, 825, 679, 533, 389, 243
11 (Albinoside IX)	109.1	$\text{C}_{50}\text{H}_{82}\text{O}_{21}$	1018.527	1041	1017	935, 853, 707, 561, 417, 271

Table 2¹H NMR and ¹³C NMR spectra data of albinosides X–XI (**1–2**) (400 MHz and 100 MHz, C₅D₅N, δ ppm, J in Hz).

Position	1		2	
	δ _H	δ _C	δ _H	δ _C
Qui-1	4.77 (d, J = 8.0 Hz)	103.2	4.75 (d, J = 8.0 Hz)	103.0
2	4.44 (dd, J = 9.0, 8.0 Hz)	77.1	4.45 (dd, J = 9.0, 8.0 Hz)	76.8
3	4.33 (dd, J = 9.0, 9.0 Hz)	79.6	4.33 (t, J = 9.0, 9.0 Hz)	79.4
4	3.63 (dd, J = 9.0, 9.0 Hz)	78.0	3.65 (dd, J = 9.0, 9.0 Hz)	77.7
5	3.67 (dq, J = 9.0, 6.0 Hz)	73.2	3.67 (dq, J = 9.0, 6.0 Hz)	72.9
6	1.59 (d, J = 6.0 Hz)	19.7	1.61 (d, J = 6.0 Hz)	19.1
Rha-1	5.89 (brs)	100.4	5.88 (brs)	100.2
2	4.79 (dd, J = 3.0, 2.0 Hz)	70.7	4.80 (dd, J = 3.0, 2.0 Hz)	70.5
3	5.76 (m [*])	79.7	5.75 (t, J = 8.8 Hz)	79.5
4	4.32 (dd, J = 9.9, 9.9 Hz)	86.1	4.33 (dd, J = 9.9, 9.9 Hz)	85.9
5	4.83 (dq, J = 9.9, 6.5 Hz)	68.9	4.84 (dq, J = 9.6, 6.5 Hz)	68.6
6	1.92 (d, J = 6.5 Hz)	19.3	1.93 (d, J = 6.5 Hz)	19.3
Rha'-1	5.64 (brs)	99.9	5.64 (brs)	99.8
2	5.88 (dd, J = 3.0, 2.0 Hz)	74.3	5.88 (dd, J = 3.0, 2.0 Hz)	73.9
3	4.15 (m [*])	81.9	4.34 (dd, J = 9.0, 3.0 Hz)	79.4
4	3.66 (dd, J = 9.9, 9.9 Hz)	75.9	3.68 (dd, J = 9.9, 9.9 Hz)	75.6
5	3.86 (dq, J = 9.9, 6.5 Hz)	72.6	3.85 (dq, J = 9.9, 6.5 Hz)	72.3
6	1.49 (d, J = 6.5 Hz)	18.8	1.49 (d, J = 6.5 Hz)	18.6
Fuc-1	5.29 (d, J = 7.6 Hz)	107.1	5.30 (d, J = 7.6 Hz)	106.9
2	4.33 (dd, J = 9.2, 7.6 Hz)	74.2	4.33 (dd, J = 9.6, 7.6 Hz)	73.7
3	4.25 (dd, J = 9.2, 3.0 Hz)	80.9	4.25 (dd, J = 9.2, 3.0 Hz)	81.7
4	5.63 (brs)	74.9	5.64 (brs)	74.7
5	3.93 (q, J = 6.0 Hz)	70.9	3.93 (q, J = 6.0 Hz)	70.7
6	1.32 (d, J = 6.4 Hz)	17.7	1.31 (d, J = 6.4 Hz)	17.4
conv-1		173.2		
2a	2.82 (ddd, J = 10.4, 7.6, 2.0 Hz)	35.2		
2b	2.43 (ddd, J = 10.4, 7.6, 2.0 Hz)			
11	3.79–3.82 (m [*])	81.7		
14	0.90 (t, J = 7.2 Hz)	14.8		
jal-1				173.0
2a			2.88 (ddd, J = 16.4, 10.4, 6.0 Hz)	35.6
2b			2.43 (ddd, J = 16.4, 10.4, 6.0 Hz)	
11			3.82–3.88 (m [*])	80.9
16			0.85 (t, J = 6.8 Hz)	14.7
tga-1		168.2		168.0
2		129.7		129.4
3	6.99 (dq, J = 7.2, 1.6 Hz)	138.3	6.99 (dq, J = 7.2, 1.6 Hz)	138.0
4	1.49 (d, J = 7.2 Hz)	14.9	1.49 (d, J = 7.2 Hz)	14.6
5	1.85 (brs)	13.1	1.81 (brs)	12.7
tga'-1		169.0		169.0
2		129.6		129.3
3	7.13 (dq, J = 7.2, 1.2 Hz)	138.8	7.13 (dq, J = 7.2, 1.2 Hz)	138.5
4	1.47 (d, J = 7.2 Hz)	15.1	1.45 (d, J = 7.2 Hz)	14.5
5	1.87 (brs)	13.0	1.86 (brs)	12.7

Chemical shifts δ are in ppm relative to TMS.

Chemical shifts marked with an asterisk (*) indicate overlapped signals.

Abbreviations: qui, quinovose; rha, rhamnose; fuc, fucose; conv, 11-hydroxytetradecanoyl; jal, 11-hydroxyhexadecanoyl; tga, tigloyl.

δ_H 4.32; **2**: δ_H 4.33) with C-1 of fucose (**1**: δ_C 107.1; **2**: δ_C 106.9); and (d) H-3 of fucose (δ_H 4.25) with C-1 of Rha' (**1**: δ_C 99.9; **2**: δ_C 99.8). Accordingly, the glycosidic acid for compound **1**, named as albinosidic acid H (**12**), corresponded to (11S)-hydroxytetradecanoic acid 11-O-α-L-rhamnopyranosyl-(1→3)-O-β-D-fucopyranosyl-(1→4)-O-α-L-rhamnopyranosyl-(1→2)-O-6-deoxy-β-D-glucopyranoside, while the structure for albinosidic acid I (**13**), the glycosidic acid for compound **2**, was characterized as (11S)-hydroxyhexadecanoic

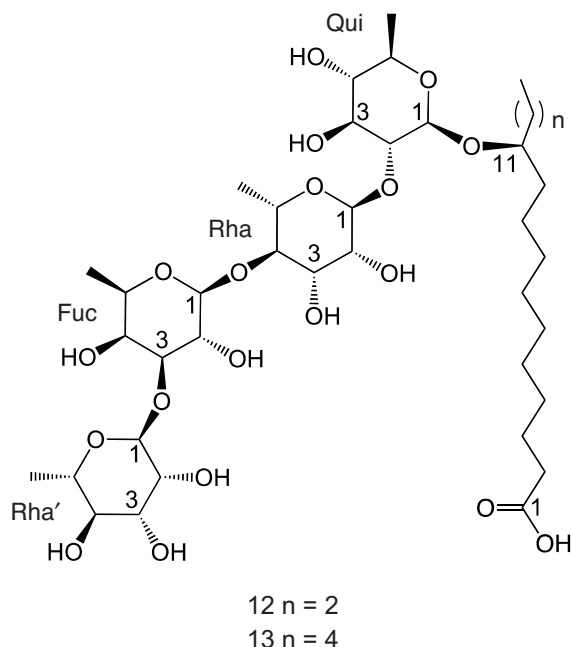
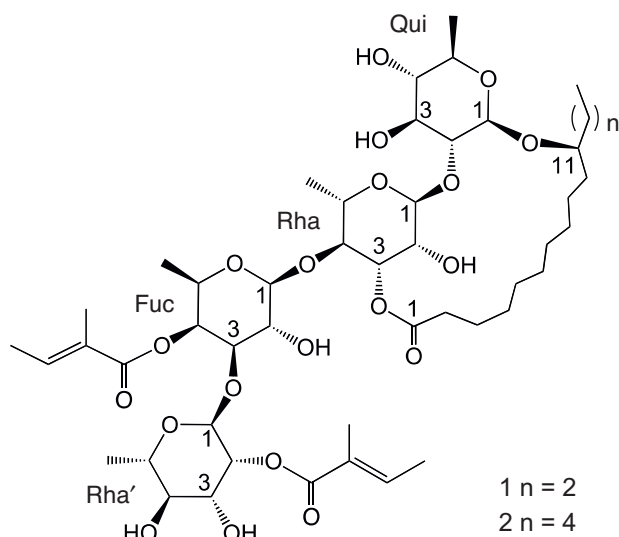
acid 11-O-α-L-rhamnopyranosyl-(1→3)-O-β-D-fucopyranosyl-(1→4)-O-α-L-rhamnopyranosyl-(1→2)-O-6-deoxy-β-D-glucopyranoside. For the natural products, the macrolactonization site was located by the ³J_{CH} correlations between H-3 (**1**: δ_H 5.76; **2**: δ_H 5.75) of Rha with C-1 (**1**: δ_C 173.2; **2**: δ_C 173.0) of the fatty acid in the HMBC spectrum. The tigloyl group linkages were at the same location on the oligosaccharide core for **1** and **2**. Thus, H-2 (δ_H 5.88) of Rha' correlated with the C-1 (**1**: δ_C 168.2; **2**: δ_C 168.0) signal

Table 3Modulation of vinblastine cytotoxicity in drug sensitive MCF-7 and multidrug-resistant MCF-7/Vin by **1** and **2**.

Compound ^a	IC ₅₀ (μg/ml)			Reversal fold ^c		
	MCF-7/Vin ⁻	MCF-7/Vin ⁺	MCF-7 sens	RF _{MCF-7/Vin⁻}	RF _{MCF-7/Vin⁺}	RF _{MCF-7 sens}
Vinblastine	1.08 ± 0.06	1.37 ± 0.23	0.047 ± 0.01			
1	0.98 ± 0.16	0.79 ± 0.098	0.015 ± 0.02	1.1	1.7	3.1
2	<0.00064	<0.00064	<0.00064	>1687.5	>2140.6	>73.4
Reserpine ^b	0.037 ± 0.01	0.31 ± 0.19	0.003 ± 0.001	29.2	4.4	15.7

^a Serial dilutions from 0.00064 to 10 μg/ml of vinblastine in the presence or absence of glycolipid (25 μg/ml).^b Reserpine = 5 μg/ml as positive control.^c RF = IC₅₀ Vinblastine/IC₅₀ Vinblastine in the presence of glycolipid. Each value represents the mean ± SD from three independent experiments.

of one tigloyl unit (tga) and H-4 (**1**: δ_{H} 5.63; **2**: δ_{H} 5.64) of Fuc correlated with C-1 (δ_{C} 169.0) of the second tigloyl group (tga').



Modulation of multidrug resistance

Based on our previous results (Cruz-Morales et al., 2012, 2016), a modulation assay was used to identify chemosensitizers through the potentiation of vinblastine susceptibility by compounds **1** and **2** in MDR cells (MCF-7/Vin⁺ cells), using the cytotoxicity screening with SRB. The reversal fold values (RF_{MCF-7/Vin⁺}) are included in Table 3. A moderate activity was displayed by the non-cytotoxic albinoside X (**1**) (RF 2.0) which was similar to that reported for albinosides I (**3**, RF 3.1), II (**4**, RF 2.6), V (**7**, RF 2.3), and VI (**8**, RF 2.1) (Cruz-Morales et al., 2012, 2016). Albinoside XI (**2**) was moderately active against various carcinoma cell lines, including the MCF-7/Vin⁺ cells (IC₅₀ 4–9 μM). Consequently, its strong reversal activity (RF >2140.6) could be the result of an additive synergism between cytotoxicity and modulation of efflux pumps.

Conclusion

For the HPLC-MS profiling, low resolution FABMS in negative mode was used for recording the fragmentation pattern of each eluted fraction and pure compounds, since this is the only procedure among the soft ionization techniques that provides a clear sequence for the fragmentation along the oligosaccharide core. Peaks resulting from the anomeric cleavages in conjunction with those produced by eliminations of the esterifying residues are abundant. In consequence, the intensity for some of the deprotonated molecule ions [M-H]⁻ was low. However, ESIMS provided highly abundant [M-H]⁻ ions in negative mode, as well as adduct ion [M+Na]⁺ detected in positive mode. Thus, this profiling based on HPLC-MS allowed the identification of novel compounds **1** and **2** as well as known compounds **3–11**.

The unexpected susceptibility of the MDR MCF-7/Vin⁺ cells to compound **2** could represent an example of collateral hypersensitivity, i.e., the MCF-7 strain, in adapting to vinblastine also acquired hypersensitivity to alternative cytotoxins as albinosides. Therefore, the additive synergism expressed in the reversal activity of **2** could be significant in therapy regimes with combination of anti-cancer drugs and deserves further studies. These results support the potential of resin glycosides as inhibitors of multidrug efflux pumps in mammalian cancer cells, where glycoprotein-P is the prevailing translocase responsible for the resistant phenotype.

Authors' contributions

JCG and SCM performed the isolation of novel compounds and completed their structure elucidation. DRR prepared derivatives and accomplished HPLC-MS dereplication. MFS performed the bioassays and supervised HPLC procedures. RPM planned the study and contributed to the preparation of the document final version.

Ethical disclosures

Protection of human and animal subjects. The authors declare that no experiments were performed on humans or animals for this study.

Confidentiality of data. The authors declare that no patient data appear in this article.

Right to privacy and informed consent. The authors declare that no patient data appear in this article.

Conflicts of interest

The authors declare no conflicts of interest.

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