

Supporting Information

Psiguadials A and B, Two Novel Meroterpenoids with Unusual Skeletons from the Leaves of *Psidium guajava*

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General Experimental Procedures

All melting points were obtained on an X-5 micro melting point apparatus without correction. Optical rotations were measured on a Jasco P-1020 polarimeter with a 2 mm cell at room temperature. UV spectra were determined on a Jasco V-550 UV/VIS spectrophotometer. IR spectra were recorded on a Jasco FT/IR-480 plus Fourier Transform infrared spectrometer using KBr pellets. ESI-MS spectra were carried out on a Finnigan LCQ Advantage Max ion trap mass spectrometer. HR-EI-MS spectra were acquired on Thermo MAT 95 XP mass spectrometer. NMR spectra were measured on Bruker AV-500 (^1H : 500 MHz, ^{13}C : 125 MHz) spectrometer with TMS as internal standard. TLC analyses were carried out using precoated silica gel GF₂₅₄ plates (Qingdao Marine Chemical Plant, Qingdao, P. R. China). Column chromatography was performed on Silica gel (200-300 mesh, Qingdao Marine Chemical Plant, Qingdao, P. R. China), reversed-phase C₁₈ silica gel (Merck, Darmstadt, Germany) and Sephadex LH-20 (Pharmacia Biotec AB, Uppsala, Sweden). All solvents used in column chromatography and HPLC were of analytical grade (Shanghai Chemical Plant, Shanghai, P. R. China) and chromatographic grade (Fisher Scientific, New Jersey, U. S. A), respectively.

Plant material

The air-dried leaves of *Psidium guajava* Linn. were collected from Conghua county, Guangdong Province, P. R. China, in October of 2007, and authenticated by Prof. Shu-Yuan Li (Guangdong Pharmaceutical University). A voucher specimen (No.

2007101801) was deposited in the Institute of Traditional Chinese Medicine & Natural Products, Jinan University, Guangzhou, P. R. China.

Extraction and Isolation

The air-dried powdered material (20.0 kg) was percolated with 95 % EtOH for three times at room temperature. After removing the solvent, the alcohol extract (1800.0 g) was suspended in H₂O and extracted by petroleum ether (PE), CHCl₃ and *n*-butanol successively to obtain the PE extract (300.0 g), CHCl₃ extract (500.0 g) and *n*-butanol extract (700.0 g), respectively. The PE extract was subjected to a silica gel column (200-300 mesh) eluted with PE/EtOAc (100/0 to 1/1; v/v) to obtain nine fractions P1-P9. Fraction P2 (21.0 g) was chromatographed over silica gel column to yield seven subfractions P2a-P2g using PE/EtOAc (100/0 to 100/10; v/v) as solvent system. Subfraction P2b was further separated by silica gel column using PE/EtOAc (100/0 to 100/10; v/v) as solvent system to afford compounds **1** (30 mg) and **2** (25 mg). Subfraction P2d was further chromatographed over silica gel column using the same solvent system to give compounds **3** (20 mg) and **4** (15 mg), respectively.

Cell lines and cell culture

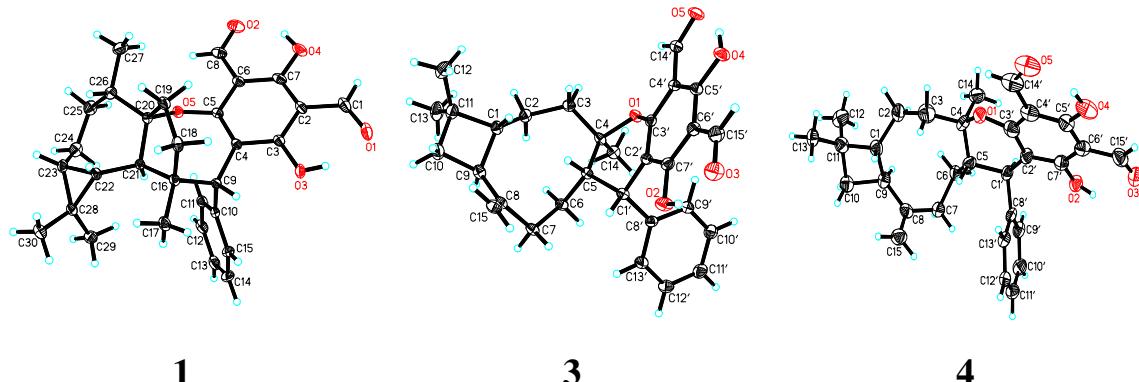
Human hepatoma carcinoma cell line HepG2 was obtained from the American Type Culture Collection. P-glycoprotein-overexpressing drug-resistant cell line HepG2/ADM was kindly provided by Prof. Kwok-Pui Fung (Department of Biochemistry, The Chinese

University of Hong Kong). Both cell lines were cultured in the RPMI 1640 medium, supplemented with 10% NBS (v/v) at 37 °C in a humidified atmosphere of 5% CO₂ (v/v). HepG2/ADM cell line was maintained in medium with 1.2 µM doxorubicin

Cell viability assay

Cells were cultured in 96-well plates for 24 h. After that, cells were treated with compounds **1-4** at various concentrations for 72 h. 30 µL of MTT solution (5 mg/mL) was added into each well and incubated for another 4 h. Subsequently, medium was discarded and 100 µL of DMSO was added to dissolve the produced formazan. The absorbance was measured at 570 nm using a microplate Reader (Thermo scientific multiskan MK3, USA). IC₅₀ values were calculated from cell survival curves using Prism software.

Single crystal X-ray data and structures of 1, 3 and 4.



X-ray data for compound 1

Identification code	pg24
Empirical formula	C ₃₀ H ₃₄ O ₅
Formula weight	474.57
Temperature	293(2) K
Wavelength	1.54178 Å
Crystal system, space group	Orthorhombic, P2(1)2(1)2(1)
Unit cell dimensions	a = 11.1759(2) Å alpha = 90 deg. b = 13.5506(2) Å beta = 90 deg. c = 15.9913(2) Å gamma = 90 deg.
Volume	2421.73(6) Å ³
Z, Calculated density	4, 1.302 Mg/m ³
Absorption coefficient	0.700 mm ⁻¹
F(000)	1016
Crystal size	0.42 x 0.40 x 0.39 mm
Theta range for data collection	4.28 to 63.21 deg.
Limiting indices	-12<=h<=12, -14<=k<=15, -17<=l<=18
Reflections collected / unique	11264 / 3885 [R(int) = 0.0287]
Completeness to theta = 63.21	99.1 %
Max. and min. transmission	0.7719 and 0.7575
Refinement method	Full-matrix least-squares on F ²
Data / restraints / parameters	3885 / 0 / 319
Goodness-of-fit on F ²	1.086
Final R indices [I>2sigma(I)]	R1 = 0.0405, wR2 = 0.1124
R indices (all data)	R1 = 0.0427, wR2 = 0.1136
Absolute structure parameter	0.1(2)
Largest diff. peak and hole	0.274 and -0.278 e.Å ⁻³

X-ray data for compound **3**

Identification code	pg19
Empirical formula	C30 H34 O5
Formula weight	474.57
Temperature	150(2) K
Wavelength	1.54178 Å
Crystal system, space group	Tetragonal, P4(1)2(1)2
Unit cell dimensions	a = 11.14980(10) Å alpha = 90 deg. b = 11.14980(10) Å beta = 90 deg. c = 40.4482(8) Å gamma = 90 deg.
Volume	5028.44(12) Å ³
Z, Calculated density	8, 1.254 Mg/m ³
Absorption coefficient	0.674 mm ⁻¹
F(000)	2032
Crystal size	0.24 x 0.15 x 0.15 mm
Theta range for data collection	4.11 to 62.50 deg.
Limiting indices	-12<=h<=12, -12<=k<=11, -44<=l<=46
Reflections collected / unique	8734 / 3928 [R(int) = 0.0256]
Completeness to theta = 62.50	98.8 %
Max. and min. transmission	0.9056 and 0.8549
Refinement method	Full-matrix least-squares on F ²
Data / restraints / parameters	3928 / 0 / 317
Goodness-of-fit on F ²	0.964
Final R indices [I>2sigma(I)]	R1 = 0.0312, wR2 = 0.0687
R indices (all data)	R1 = 0.0386, wR2 = 0.0705
Absolute structure parameter	0.1(2)
Largest diff. peak and hole	0.125 and -0.171 e. Å ⁻³

X-ray data for compound **4**

Identification code	pg8
Empirical formula	C30 H34 O5
Formula weight	474.57
Temperature	293(2) K
Wavelength	1.54178 Å
Crystal system, space group	Monoclinic, P2(1)
Unit cell dimensions	a = 6.5752(10) Å alpha = 90 deg. b = 17.504(3) Å beta = 90.862(16) deg. c = 10.7096(18) Å gamma = 90 deg.
Volume	1232.4(3) Å ³
Z, Calculated density	2, 1.279 Mg/m ³
Absorption coefficient	0.688 mm ⁻¹
F(000)	508
Crystal size	0.35 x 0.30 x 0.28 mm
Theta range for data collection	4.13 to 59.97 deg.
Limiting indices	-7<=h<=4, -19<=k<=19, -10<=l<=12
Reflections collected / unique	3122 / 2574 [R(int) = 0.0556]
Completeness to theta = 59.97	93.6 %
Refinement method	Full-matrix least-squares on F ²
Data / restraints / parameters	2574 / 1 / 316
Goodness-of-fit on F ²	1.472
Final R indices [I>2sigma(I)]	R1 = 0.1093, wR2 = 0.3091
R indices (all data)	R1 = 0.1220, wR2 = 0.3317
Absolute structure parameter	0.0(8)
Largest diff. peak and hole	0.762 and -0.359 e. Å ⁻³

The X-ray diffraction data of **1**, **3** and **4** were collected on an Oxford Gemini R diffractometer with CuK α radiation ($\lambda = 1.54018\text{\AA}$). The crystal structure was solved and refined using the SHELXS-97 software package ^[1]. In the structure refinements, non-hydrogen atoms were refined anisotropically. Hydrogen atoms bonded to carbons were placed on the geometrically ideal positions by the ‘ride on’ method. Hydrogen atoms bonded to oxygen were located by the difference Fourier method and were included in the calculation of structure factors with isotropic temperature factors.

Complete lists of refined atomic coordinates and relevant information in standard CIF format for the structures reported in this paper have been deposited with the Cambridge Crystallographic Data Center (CCDC no. 774761-774763). These materials are available free of charge via application to CCDC, 12 Union Road, Cambridge CB2 1EZ, UK [Fax: (+44) 1223-336033; e-mail: deposit@ccdc.cam.ac.uk].

Reference

- [1] G. M. Sheldrick SHELXTL, version 5.1: Structure Determination Software Programs Package, Bruker Analytical X-ray Systems, Inc., Madison, WI, USA, 1997.

Quantum chemical CD data for 1-4

Coordinates and optimized energies of computations (method: B3LYP/6-31G)

Cartesian coordinate of compound 1 optimized:

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-1.701727	-2.855290	-0.138689
2	1	0	-2.505823	-2.936704	-0.879589
3	6	0	-0.848613	-4.129197	-0.273146
4	1	0	-1.484673	-5.014786	-0.162589
5	1	0	-0.354881	-4.199610	-1.247639
6	1	0	-0.076183	-4.185852	0.498686
7	6	0	-0.939009	-1.569753	-0.550476
8	6	0	-0.425655	-1.643641	-2.015664
9	1	0	-1.021728	-2.378692	-2.565558
10	1	0	0.608126	-1.997753	-2.049980
11	6	0	-0.598286	-0.235834	-2.627402
12	1	0	-1.382877	-0.250165	-3.392580
13	1	0	0.312857	0.121884	-3.119289
14	6	0	-1.025024	0.704451	-1.466793
15	6	0	-1.906086	1.849981	-1.991839
16	1	0	-2.799987	1.461932	-2.488434
17	1	0	-2.230452	2.520737	-1.191600
18	1	0	-1.354335	2.449793	-2.726480
19	6	0	0.267911	1.334748	-0.827179
20	1	0	0.646225	1.988139	-1.617249
21	6	0	0.006890	2.258723	0.371309
22	6	0	-0.093440	1.784959	1.689156
23	1	0	0.034733	0.726617	1.894452
24	6	0	-0.334546	2.657482	2.751851
25	1	0	-0.406955	2.264185	3.762690
26	6	0	-0.467643	4.027853	2.522837
27	1	0	-0.648807	4.707484	3.351183
28	6	0	-0.349015	4.517427	1.222004
29	1	0	-0.431000	5.584129	1.029838
30	6	0	-0.111019	3.640784	0.162496
31	1	0	-0.002747	4.036485	-0.844098

32	6	0	1.433051	0.395022	-0.516591
33	6	0	2.735119	0.902370	-0.765560
34	8	0	2.837916	2.077436	-1.386354
35	1	0	3.817657	2.280274	-1.478359
36	6	0	3.923931	0.223247	-0.373055
37	6	0	5.222486	0.793395	-0.657143
38	1	0	6.098777	0.219556	-0.321206
39	8	0	5.389552	1.872368	-1.248028
40	6	0	3.800043	-0.997028	0.325391
41	8	0	4.906971	-1.618158	0.727370
42	1	0	4.610964	-2.453401	1.202889
43	6	0	2.518344	-1.545537	0.579513
44	6	0	2.423923	-2.776975	1.329557
45	1	0	1.416486	-3.148126	1.556534
46	8	0	3.409083	-3.414398	1.741361
47	6	0	1.353314	-0.846824	0.118954
48	8	0	0.183830	-1.471465	0.404766
49	6	0	-1.766438	-0.249630	-0.483755
50	1	0	-1.697657	0.145975	0.528631
51	6	0	-3.239130	-0.487606	-0.802651
52	1	0	-3.438215	-0.744394	-1.843902
53	6	0	-4.390420	0.223892	-0.096779
54	6	0	-4.168514	1.334369	0.918519
55	1	0	-4.113282	2.307380	0.414974
56	1	0	-5.011270	1.376216	1.620664
57	1	0	-3.255394	1.225490	1.507601
58	6	0	-5.634065	0.462023	-0.942637
59	1	0	-5.801458	-0.352477	-1.656883
60	1	0	-6.528547	0.539193	-0.310843
61	1	0	-5.550451	1.396121	-1.513409
62	6	0	-4.078984	-1.226450	0.228246
63	1	0	-4.789039	-1.950889	-0.173040
64	6	0	-3.394518	-1.688425	1.504329
65	1	0	-2.888432	-0.848516	1.987486
66	1	0	-4.145415	-2.030471	2.228070
67	6	0	-2.373104	-2.829169	1.253304
68	1	0	-2.882406	-3.795693	1.364417
69	1	0	-1.601279	-2.806507	2.033038

Cartesian coordinate of compound 2 optimized:

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	3.184757	0.259166	1.536143
2	6	0	4.052804	0.003207	2.780967
3	1	0	4.892552	0.706475	2.826544
4	1	0	3.465840	0.120543	3.699869
5	1	0	4.468327	-1.011027	2.781079
6	6	0	4.032804	0.168283	0.230766
7	1	0	3.934241	1.127099	-0.291200
8	6	0	5.539547	-0.205827	0.123428
9	1	0	6.270927	0.608085	0.185944
10	1	0	5.831784	-0.995135	0.824305
11	6	0	5.237977	-0.804481	-1.294799
12	6	0	5.459297	0.210852	-2.423360
13	1	0	5.046963	-0.156839	-3.371278
14	1	0	4.996589	1.181770	-2.214531
15	1	0	6.531732	0.384578	-2.575619
16	6	0	5.932452	-2.124009	-1.631619
17	1	0	5.795019	-2.862938	-0.833221
18	1	0	5.536573	-2.556121	-2.560151
19	1	0	7.011586	-1.978661	-1.770371
20	6	0	3.746919	-0.923833	-0.817192
21	1	0	3.667879	-1.898999	-0.315889
22	6	0	2.508657	-0.723570	-1.679334
23	1	0	2.545547	-1.357198	-2.576422
24	1	0	2.461322	0.312173	-2.040869
25	6	0	1.223092	-1.112422	-0.917724
26	1	0	1.255984	-2.197779	-0.759007
27	1	0	0.359276	-0.936718	-1.567782
28	6	0	0.906875	-0.532576	0.492867
29	8	0	-0.201034	-1.369249	0.975516
30	6	0	-1.429497	-1.201142	0.430080
31	6	0	-2.323107	-2.289682	0.607335
32	6	0	-1.879160	-3.483916	1.356427
33	1	0	-0.830464	-3.428148	1.703395
34	8	0	-2.553542	-4.468421	1.602918
35	6	0	-3.604882	-2.191925	0.017599

36	8	0	-4.469380	-3.186789	0.157693
37	1	0	-5.295317	-2.930966	-0.343374
38	6	0	-3.964733	-1.039550	-0.743273
39	6	0	-5.273259	-0.952197	-1.356924
40	1	0	-5.496680	-0.033107	-1.918921
41	8	0	-6.136647	-1.839512	-1.279233
42	6	0	-3.034727	0.020816	-0.866576
43	8	0	-3.440090	1.070281	-1.621105
44	1	0	-2.859070	1.837858	-1.469459
45	6	0	-1.770927	-0.014726	-0.257576
46	6	0	-0.790006	1.156513	-0.363983
47	1	0	-0.391773	1.201723	-1.389621
48	6	0	-1.434112	2.516746	-0.082204
49	6	0	-2.205206	2.731449	1.072308
50	1	0	-2.384015	1.905468	1.755536
51	6	0	-2.754519	3.983763	1.339498
52	1	0	-3.350075	4.129282	2.236578
53	6	0	-2.549092	5.047620	0.455894
54	1	0	-2.983447	6.021429	0.663646
55	6	0	-1.791167	4.848973	-0.696884
56	1	0	-1.627832	5.667185	-1.392867
57	6	0	-1.237001	3.592838	-0.961674
58	1	0	-0.640132	3.446009	-1.859665
59	6	0	0.396849	0.917802	0.600049
60	1	0	-0.016778	0.980592	1.617614
61	6	0	1.523895	1.956853	0.532415
62	1	0	1.104003	2.958494	0.674942
63	1	0	1.985932	1.958576	-0.462870
64	6	0	2.572511	1.681562	1.623377
65	1	0	2.095575	1.810663	2.605664
66	1	0	3.376104	2.427809	1.567708
67	6	0	2.012732	-0.763386	1.544334
68	1	0	2.388437	-1.789149	1.458586
69	1	0	1.520295	-0.702712	2.523215

Cartesian coordinate of compound 3 optimized:

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	6.066531	0.145161	-0.451935
2	1	0	6.742438	0.856689	0.035051
3	1	0	6.446501	-0.069306	-1.456630
4	6	0	4.557188	0.496648	-0.481488
5	1	0	4.312901	1.100213	0.404911
6	6	0	3.950497	1.145805	-1.702379
7	6	0	4.660927	1.909011	-2.539124
8	1	0	5.722275	2.092058	-2.392852
9	1	0	4.204038	2.381440	-3.405522
10	6	0	2.485877	0.852313	-1.927424
11	1	0	2.133422	1.349889	-2.838920
12	1	0	2.387191	-0.224749	-2.117094
13	6	0	1.571125	1.236531	-0.737187
14	1	0	2.141140	1.213559	0.195287
15	1	0	1.271446	2.279837	-0.850820
16	6	0	0.330920	0.322727	-0.608860
17	1	0	0.253928	-0.248551	-1.544631
18	6	0	-1.046836	1.070278	-0.540226
19	1	0	-1.166320	1.504779	-1.541505
20	6	0	-1.194530	2.262603	0.418396
21	6	0	-1.824105	2.165705	1.666922
22	1	0	-2.247244	1.216745	1.982536
23	6	0	-1.956659	3.276451	2.501209
24	1	0	-2.450367	3.167840	3.463465
25	6	0	-1.474838	4.521805	2.097705
26	1	0	-1.580283	5.388179	2.744981
27	6	0	-0.882776	4.647378	0.840797
28	1	0	-0.532947	5.616631	0.495160
29	6	0	-0.757688	3.532898	0.011164
30	1	0	-0.340015	3.661910	-0.984240
31	6	0	-2.137075	0.028604	-0.388155
32	6	0	-3.462619	0.327965	-0.766198
33	8	0	-3.708539	1.551812	-1.226799
34	1	0	-4.687750	1.592988	-1.443894
35	6	0	-4.519095	-0.626328	-0.682454

36	6	0	-5.859774	-0.262938	-1.080118
37	1	0	-6.632134	-1.041480	-0.989751
38	8	0	-6.171360	0.859929	-1.509249
39	6	0	-4.213595	-1.927225	-0.220316
40	8	0	-5.196256	-2.822017	-0.137020
41	1	0	-4.782340	-3.672879	0.202559
42	6	0	-2.887508	-2.269902	0.144581
43	6	0	-2.603024	-3.604575	0.614448
44	1	0	-1.563351	-3.826297	0.895621
45	8	0	-3.461683	-4.498205	0.717897
46	6	0	-1.866169	-1.270840	0.039865
47	8	0	-0.633179	-1.699197	0.364416
48	6	0	0.509185	-0.768373	0.475979
49	6	0	0.513929	-0.277453	1.928105
50	1	0	0.767136	-1.104839	2.599081
51	1	0	1.238454	0.525785	2.085528
52	1	0	-0.462995	0.105829	2.217996
53	6	0	1.703379	-1.735578	0.209883
54	1	0	1.771946	-1.929777	-0.867026
55	1	0	1.368582	-2.679567	0.652669
56	6	0	3.113569	-1.429150	0.799069
57	1	0	3.454181	-2.360923	1.269335
58	1	0	3.041278	-0.711032	1.623049
59	6	0	4.238740	-1.002062	-0.158101
60	1	0	4.140268	-1.556872	-1.101645
61	6	0	5.730226	-1.159117	0.330883
62	6	0	5.915422	-0.965420	1.842504
63	1	0	6.983185	-0.882397	2.080016
64	1	0	5.426727	-0.054880	2.208102
65	1	0	5.516325	-1.811969	2.414023
66	6	0	6.440639	-2.431322	-0.131205
67	1	0	6.351135	-2.566003	-1.215723
68	1	0	7.509936	-2.399405	0.115169
69	1	0	6.018277	-3.321619	0.353711

Cartesian coordinate of compound 4 optimized:

Standard orientation:

Center Number	Atomic Number	Atomic Type	Coordinates (Angstroms)		
			X	Y	Z
1	6	0	-5.741129	0.486129	-0.553093
2	1	0	-5.808407	0.647067	-1.634265
3	1	0	-6.511437	1.090350	-0.060845
4	6	0	-4.288615	0.638140	-0.026762
5	1	0	-4.342986	0.826095	1.052984
6	6	0	-3.282890	1.588819	-0.630609
7	6	0	-3.229209	1.869450	-1.937781
8	1	0	-3.936915	1.454756	-2.651275
9	1	0	-2.473651	2.537389	-2.343792
10	6	0	-2.308340	2.206233	0.350913
11	1	0	-1.605570	2.860567	-0.174072
12	1	0	-2.877510	2.848531	1.041005
13	6	0	-1.540897	1.181396	1.216302
14	1	0	-1.070486	1.721578	2.046803
15	1	0	-2.267773	0.516346	1.689379
16	6	0	-0.460307	0.348771	0.473592
17	1	0	-0.608002	0.464562	-0.607842
18	6	0	0.963158	0.900858	0.818736
19	1	0	1.004473	1.007739	1.913388
20	6	0	1.118901	2.309330	0.237669
21	6	0	1.094660	3.437052	1.073289
22	1	0	0.989287	3.303914	2.148089
23	6	0	1.198289	4.727859	0.544047
24	1	0	1.175898	5.585949	1.210142
25	6	0	1.329660	4.909011	-0.831255
26	1	0	1.412083	5.909685	-1.245872
27	6	0	1.365006	3.793577	-1.674466
28	1	0	1.476694	3.926529	-2.747090
29	6	0	1.263772	2.508690	-1.145991
30	1	0	1.304364	1.647470	-1.807717
31	6	0	2.081624	-0.046525	0.396867
32	6	0	3.434194	0.324311	0.442070
33	8	0	3.839777	1.541004	0.874616
34	1	0	3.082561	2.153710	0.916591
35	6	0	4.472498	-0.557655	0.049774

36	6	0	5.856454	-0.140420	0.066790
37	1	0	6.062792	0.888156	0.397647
38	8	0	6.803710	-0.869367	-0.271037
39	6	0	4.126283	-1.868439	-0.383387
40	8	0	5.085399	-2.713956	-0.767463
41	1	0	5.957422	-2.230828	-0.670390
42	6	0	2.784022	-2.299705	-0.402525
43	6	0	2.491892	-3.689542	-0.805961
44	1	0	3.401014	-4.262080	-1.068324
45	8	0	1.395347	-4.222223	-0.865299
46	6	0	1.789187	-1.371299	0.013775
47	8	0	0.536368	-1.832827	0.011877
48	6	0	-0.548079	-1.171431	0.743865
49	6	0	-0.388690	-1.551747	2.220832
50	1	0	-1.197847	-1.137948	2.830044
51	1	0	-0.403389	-2.640871	2.322843
52	1	0	0.560339	-1.188832	2.628615
53	6	0	-1.748299	-1.898427	0.079643
54	1	0	-1.455396	-2.952701	0.112247
55	1	0	-1.728561	-1.630132	-0.983244
56	6	0	-3.200815	-1.767159	0.612775
57	1	0	-3.226958	-1.439587	1.659469
58	1	0	-3.627244	-2.779232	0.622915
59	6	0	-4.119174	-0.898762	-0.254388
60	1	0	-3.825799	-1.034220	-1.304392
61	6	0	-5.682255	-1.036395	-0.208594
62	6	0	-6.238251	-1.330842	1.191466
63	1	0	-5.837422	-0.653144	1.953956
64	1	0	-7.330005	-1.222464	1.197355
65	1	0	-6.007271	-2.356514	1.504174
66	6	0	-6.293678	-1.993399	-1.231403
67	1	0	-7.388258	-1.910986	-1.245882
68	1	0	-5.928639	-1.781808	-2.243434
69	1	0	-6.046026	-3.037364	-0.997904

The complete reference:

Frisch, M. J.; Trucks, G. W.; Schlegel, H. B.; Scuseria, G. E.; Robb, M. A.; Cheeseman, J. R.; Scalmani, G.; Barone, V.; Mennucci, B.; Petersson, G. A.; Nakatsuji, H.; Caricato, M.; Li, X.; Hratchian, H. P.; Izmaylov, A. F.; Bloino, J.; Zheng, G.; Sonnenberg, J. L.; Hada, M.; Ehara, M.; Toyota, K.; Fukuda, R.; Hasegawa, J.; Ishida, M.; Nakajima, T.; Honda, Y.; Kitao, O.; Nakai, H.; Vreven, T.; Montgomery, Jr., J. A.; Peralta, J. E.;

Ogliaro, F.; Bearpark, M.; Heyd, J. J.; Brothers, E.; Kudin, K. N.; Staroverov, V. N.; Kobayashi, R.; Normand, J.; Raghavachari, K.; Rendell, A.; Burant, J. C.; Iyengar, S. S.; Tomasi, J.; Cossi, M.; Rega, N.; Millam, N. J.; Klene, M.; Knox, J. E.; Cross, J. B.; Bakken, V.; Adamo, C.; Jaramillo, J.; Gomperts, R.; Stratmann, R. E.; Yazyev, O.; Austin, A. J.; Cammi, R.; Pomelli, C.; Ochterski, J. W.; Martin, R. L.; Morokuma, K.; Zakrzewski, V. G.; Voth, G. A.; Salvador, P.; Dannenberg, J. J.; Dapprich, S.; Daniels, A. D.; Farkas, Ö.; Foresman, J. B.; Ortiz, J. V.; Cioslowski, J.; Fox, D. J. *Gaussian 09W, Revision A.02*, Gaussian, Inc., Wallingford CT, 2009.

Spectral data of 1 and 2

Psiguadials A (**1**): colorless needles (CHCl₃); [α]_D²⁵ -62.5 (c 0.15, CHCl₃); IR (KBr) ν_{max} 2964, 2933, 1632, 1428, 1308, 1172, 869, 766, 610 cm⁻¹; UV (CHCl₃) λ_{max} (log ε) 229 nm (3.37), 296 nm (3.85), 347 nm (3.33); HR-EI-MS *m/z* 474.2403 [M]⁺ (calcd for C₃₀H₃₄O₅ 474.2406).

1D and 2D NMR data of **1** (CDCl₃)

no.	δ_{H}	δ_{C}	¹ H- ¹ H COSY	HMBC	ROESY
1	-	104.1			
2	a 2.01 b 1.88	36.6	H ₂ -3	C-3, C-4	
3	a 1.90 b 1.63	37.9		C-1, C-4, C-15, C-1'	
4	-	47.8			
5	2.36 (d, 11.9)	48.7		C-2, C-3, C-4, C-6, C-11, C-1'	H-8α, H ₃ -12, H-9', H-10', H-11', H-12', H-13'
6	0.29 (dd, 8.8, 11.9)	26.3	H-5, H-7	C-4, C-7, C-11, C-12	H ₂ -2, H-3a, H-7, H-10, H ₃ -13, H ₃ -15
7	0.58	23.9		C-6, C-12	H ₃ -12, H-8β, H-10
8	α 1.26 (m) β 1.83 (m)	20.3	H-7, H-9	C-6, C-7, C-9, C-10	
9	α 1.74 (m) β 1.93 (m)	30.3		C-7, C-8, C-10, C-14	
10	2.49 (m)	39.9	H ₂ -9, H-14	C-9, C-14	H ₂ -2
11	-	19.2			
12	0.91 (s)	28.4		C-6, C-7, C-11, C-13	
13	0.24 (s)	14.3		C-7, C-11, C-12	H-6, H-8α, H ₃ -12, H ₃ -15, H-10', H-12'
14	1.09 (d, 6.8)	17.4		C-1, C-9, C-10	H ₂ -9, H-10, H-14'
15	1.13 (s)	24.9		C-3, C-4, C-1'	H-3a
1'	4.49 (s)	53.2		C-3, C-4, C-15, C-2', C-3', C-7', C-8', C-9', C-13'	H-3b, H ₃ -15
2'	-	114.2			
3'	-	166.7			
4'	-	108.2			
5'	-	168.2			
6'	-	105.0			

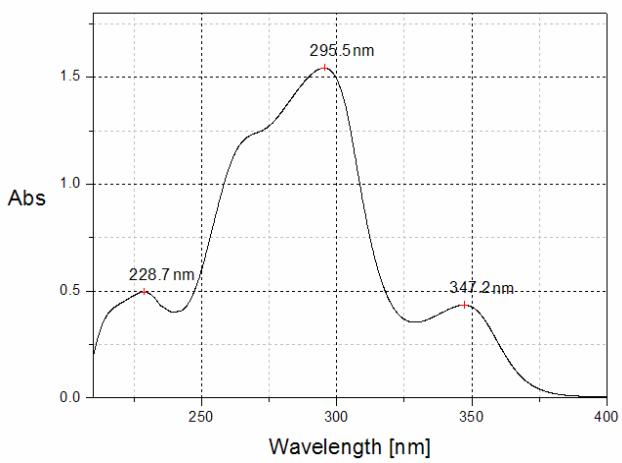
7'	-	170.5			
8'	-	140.9			
9'	7.13	130.3	C-1', C-11', C-13'		
10'	7.22	127.6	H ₂ -9', H-11'	C-8', C-12'	
11'	7.11	126.4		C-9', C-13'	
12'	7.22	127.6	H-11', H-13'	C-8', C-10'	
13'	7.13	130.3		C-1', C-9', C-11'	
14'	10.38 (s)	193.7		C-4', C-5', C-6'	
15'	10.12 (s)	191.8		C-2', C-6', C-7'	
5'-OH	13.76 (s)			C-4', C-5', C-6', C-14'	H-14', H-15'
7'-OH	13.53 (s)			C-2', C-6', C-7', C-15'	H-1', H-15'

Psiguadials B (**2**): colorless needles (CHCl₃); [α]_D²⁵ +70.0 (c 0.15, CHCl₃); IR (KBr) ν_{max} 2952, 2869, 1625, 1442, 1302, 1185, 1154, 828, 700, 607, 566 cm⁻¹; UV (CHCl₃) λ_{max} (log ε) 219 nm (3.32), 277 nm (3.69), 339 nm (3.09); HR-EI-MS *m/z* 474.2403 [M]⁺ (calcd for C₃₀H₃₄O₅ 474.2406).

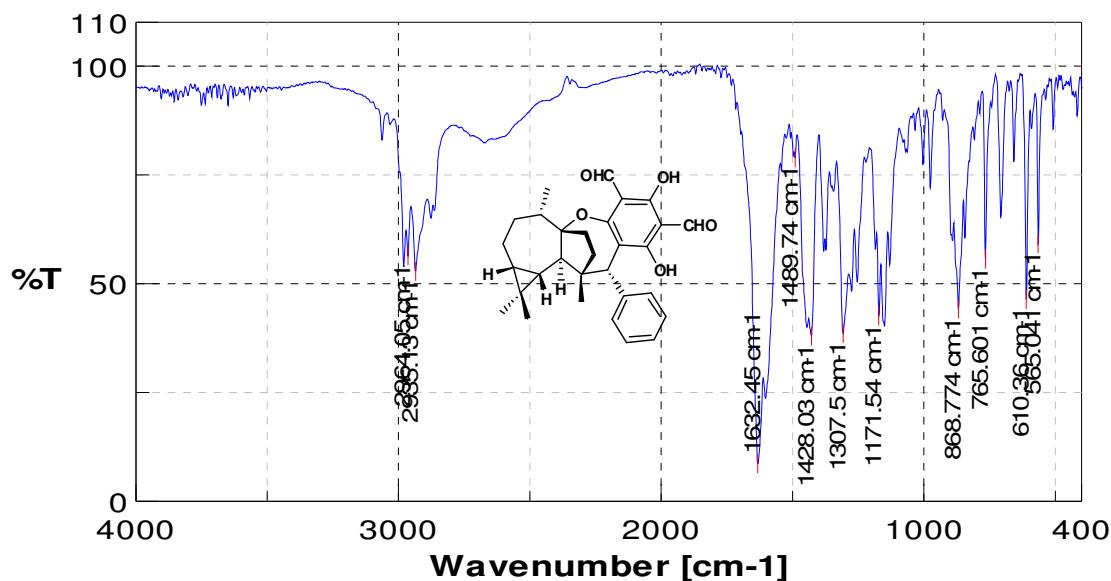
1D and 2D NMR data of **2** (CDCl₃)

no.	δ _H	δ _C	¹ H- ¹ H COSY	HMBC	ROESY
1	-	33.4			
2	2.16	37.0	H ₂ -3, H-5	C-3, C-5, C-6, C-15	H-3 α , H-11a, H ₃ -13
3	α 1.52 β 1.37	35.5		C-1, C-2, C-4, C-5, C-13, C-14	
4	-	35.1			
5	1.82 (m)	44.1		C-6	H-6 β , H-12a, H ₃ -14
6	α 1.65 (m) β 1.41 (m)	20.1	H-5, H ₂ -7		
7	α 1.93 β 1.58	29.4		C-6	
8	-	84.1			
9	1.68	50.0	H-10, H-1	C-7, C-10, C-1'	H-12b
10	1.49 (m)	23.9	H ₂ -9, H ₂ -11	C-1	H-1'
11	α 1.41 β 1.10	37.6			
12	a 2.08 b 1.29,	47.5		C-1, C-2, C-7, C-8, C-9, C-11	
13	1.02 (s)	30.6		C-3, C-14	

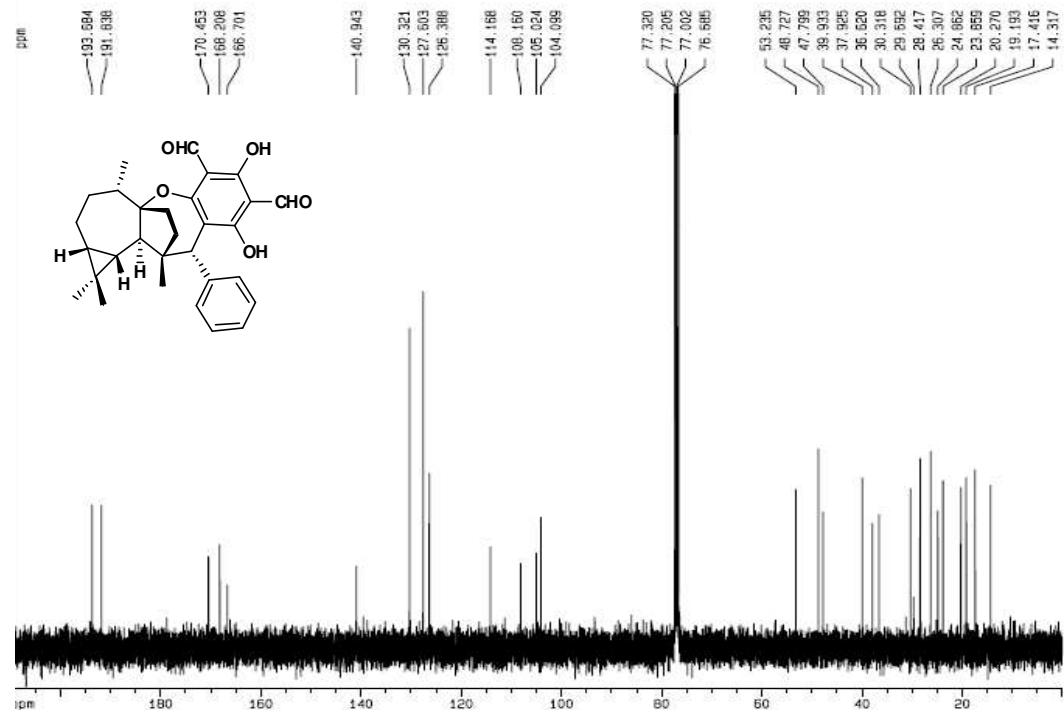
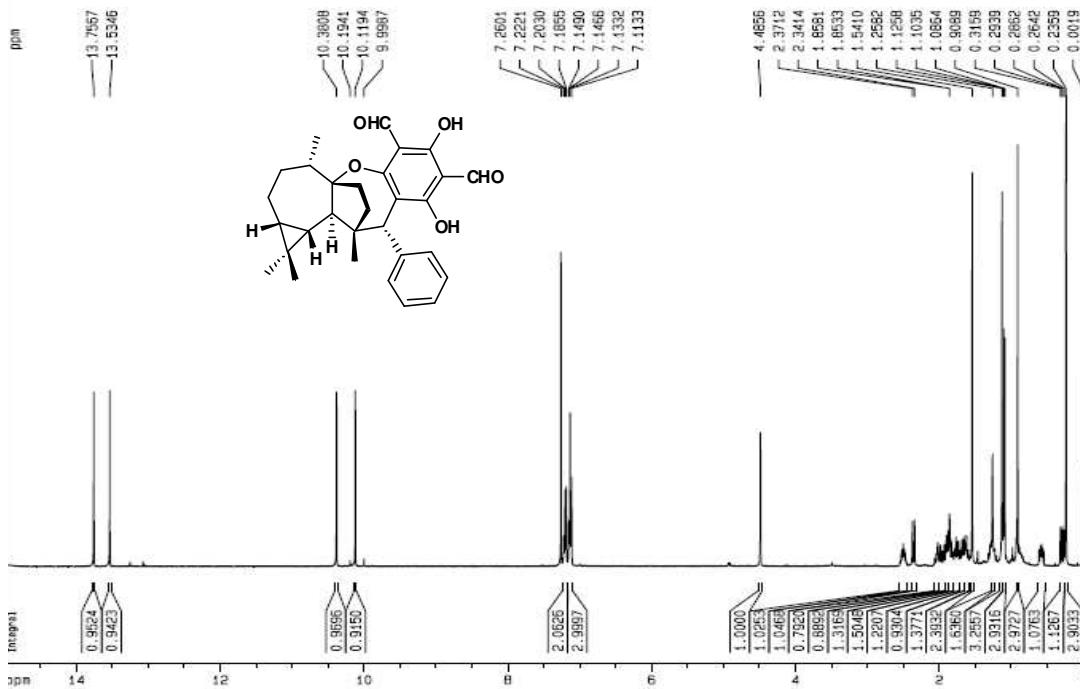
14	1.01 (s)	20.7	C-3, C-5, C-13	H-12a
15	0.86 (s)	26.1	C-1, C-11, C-12	H-3 β , H-12a
1'	3.49 (d,11.5)	40.4	C-9, C-10, C-3', C-8', C-9', C-13'	H-6 α , H-7 α
2'	-	105.7		
3'	-	163.5		
4'	-	104.6		
5'	-	168.5		
6'	-	104.2		
7'	-	169.6		
8'	-	143.4		
9'	7.18	128.2	C-1', C-11', C-13'	
10'	7.23	126.2	H ₂ -9', H-11'	C-8', C-12'
11'	7.18	128.2		C-9', C-13'
12'	7.23	126.2	H-11', H-13'	C-8', C-10'
13'	7.18	128.2		C-1', C-9', C-11'
14'	10.08 (s)	191.4		C-4', C-5', C-6'
15'	10.08 (s)	192.3		C-2', C-6', C-7'
5'-OH	13.51 (s)		C-4', C-5', C-6', C-14'	H-14', H-15'
7'-OH	13.04 (s)		C-2', C-6', C-7', C-15'	H-1', H-15'

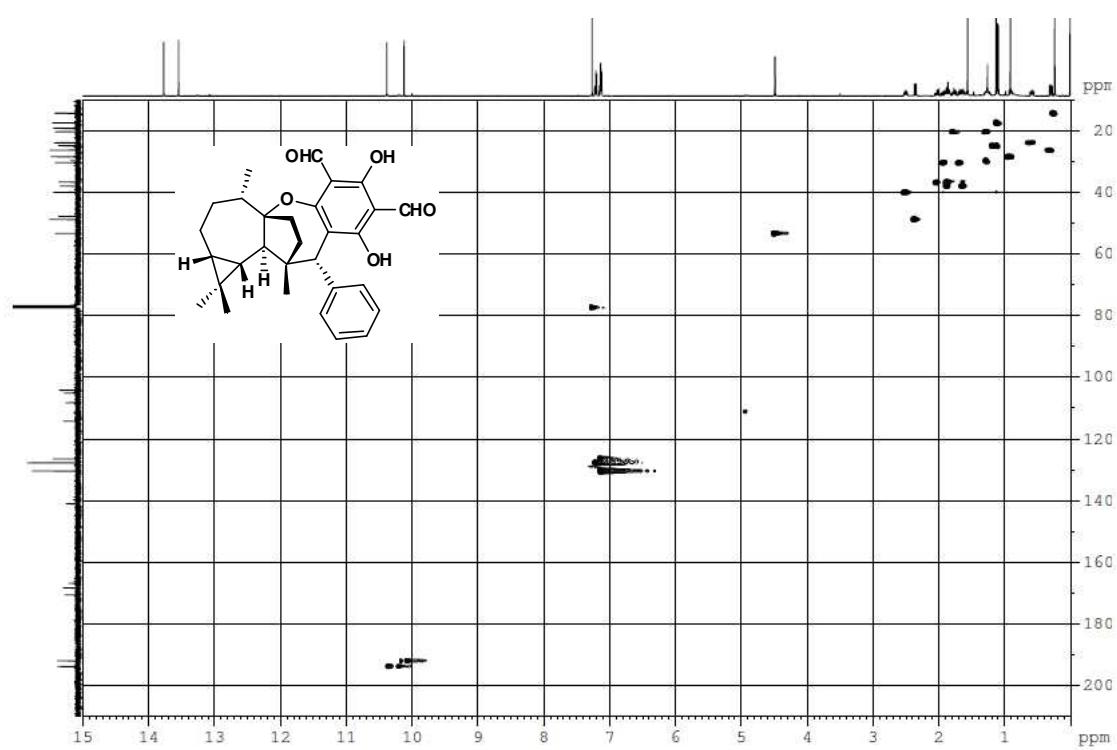
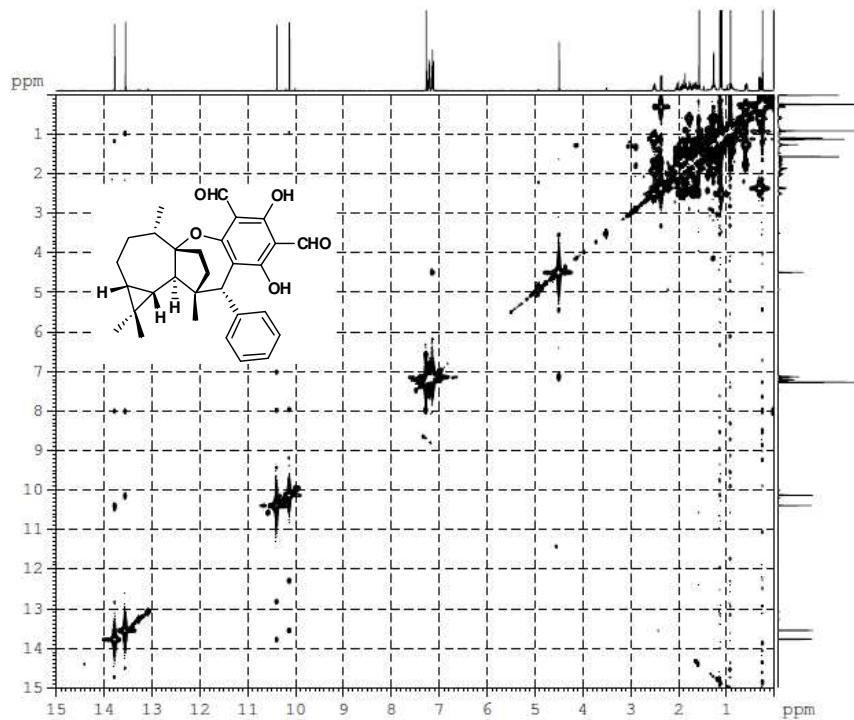


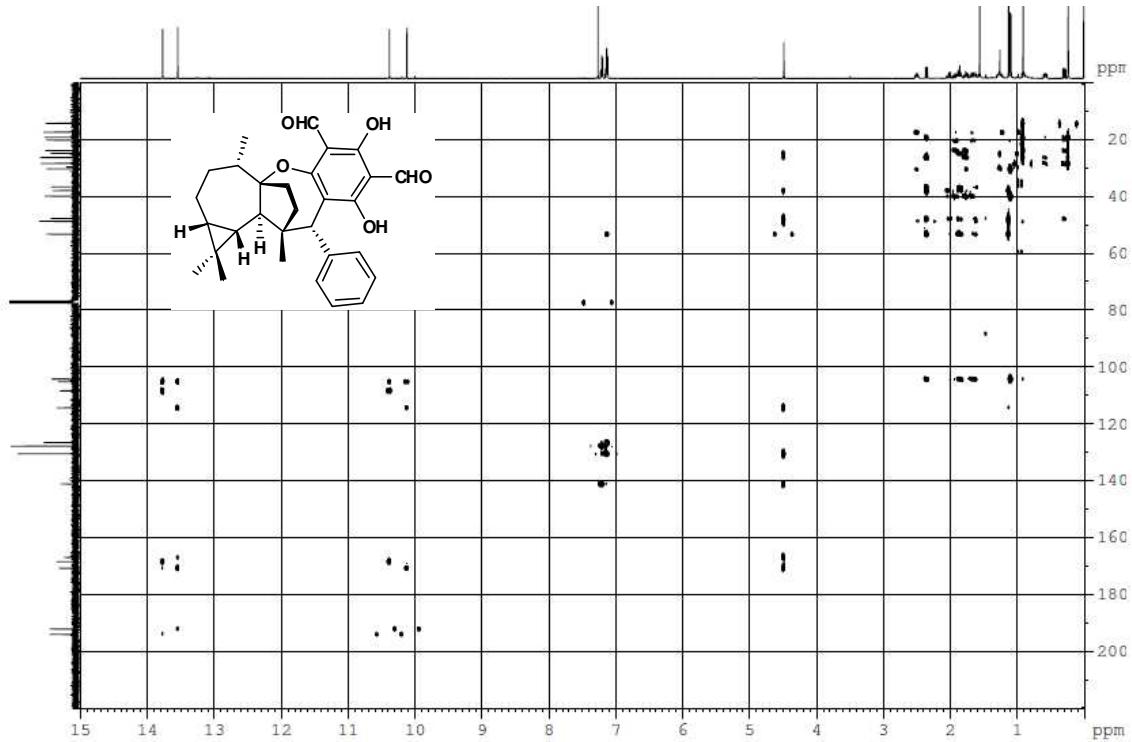
UV spectrum of **1** (CHCl_3)



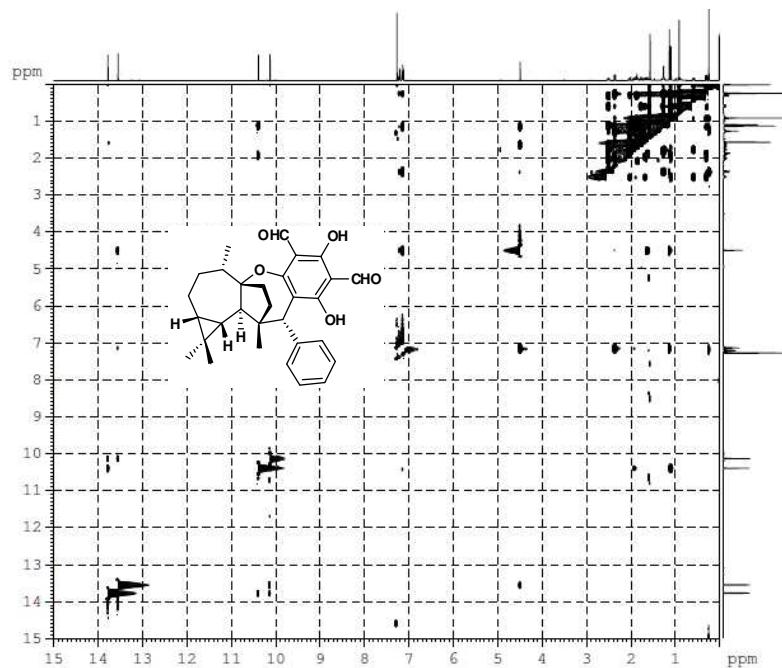
IR spectrum of **1** (KBr disc)



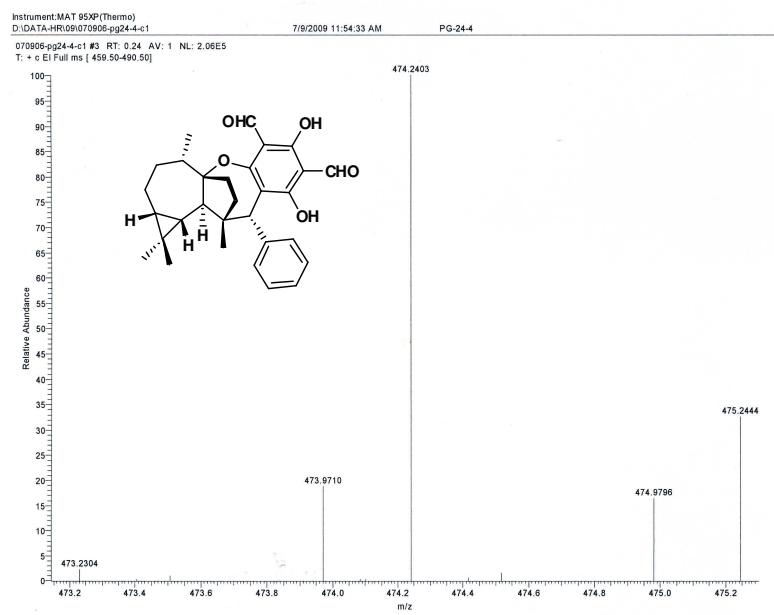




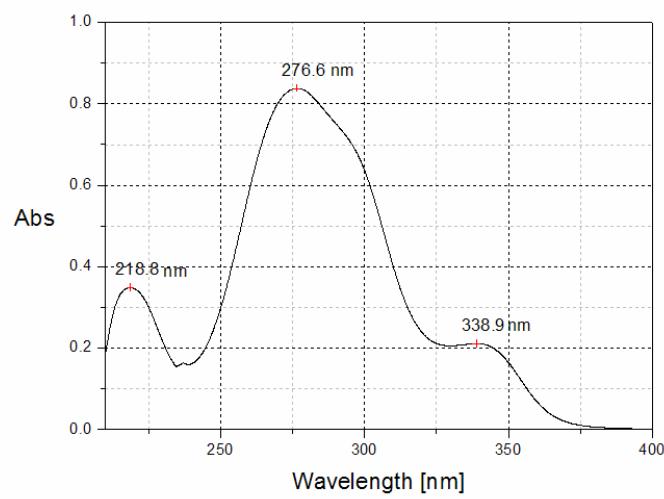
HMBC spectrum of **1** in CDCl_3



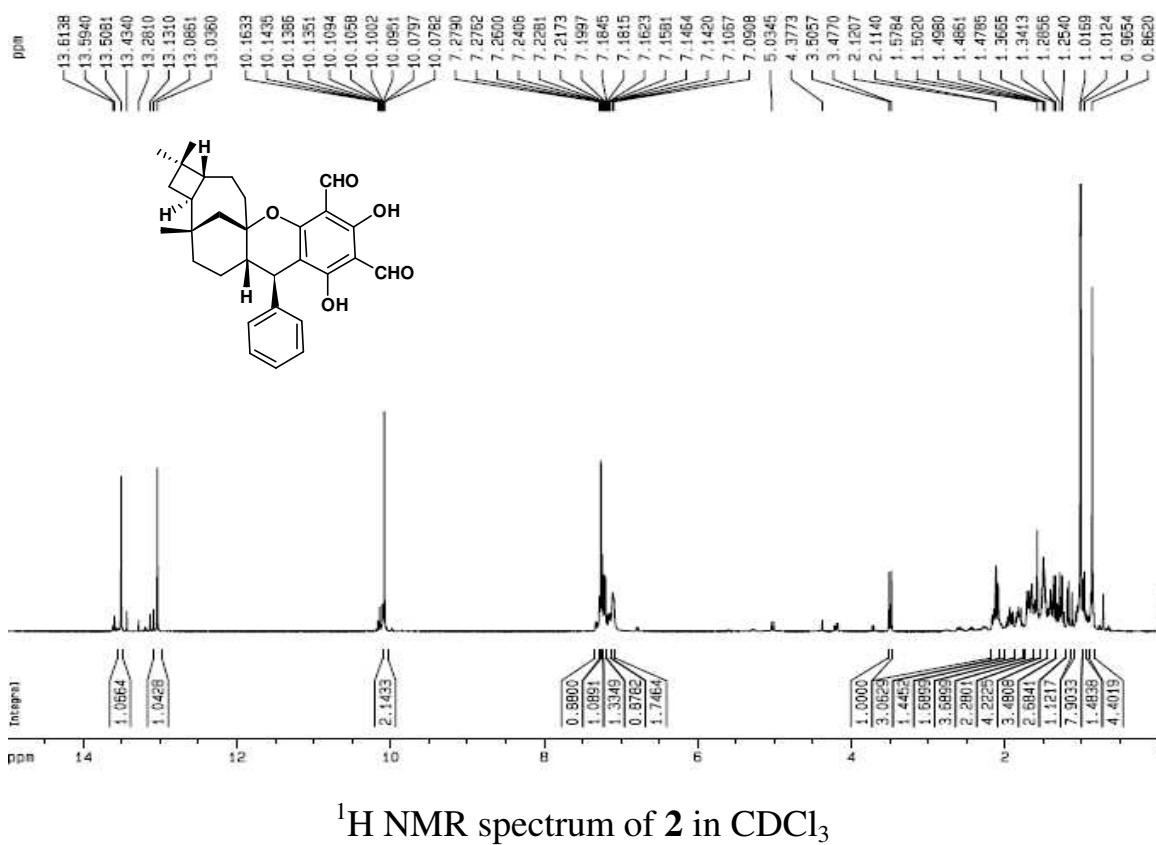
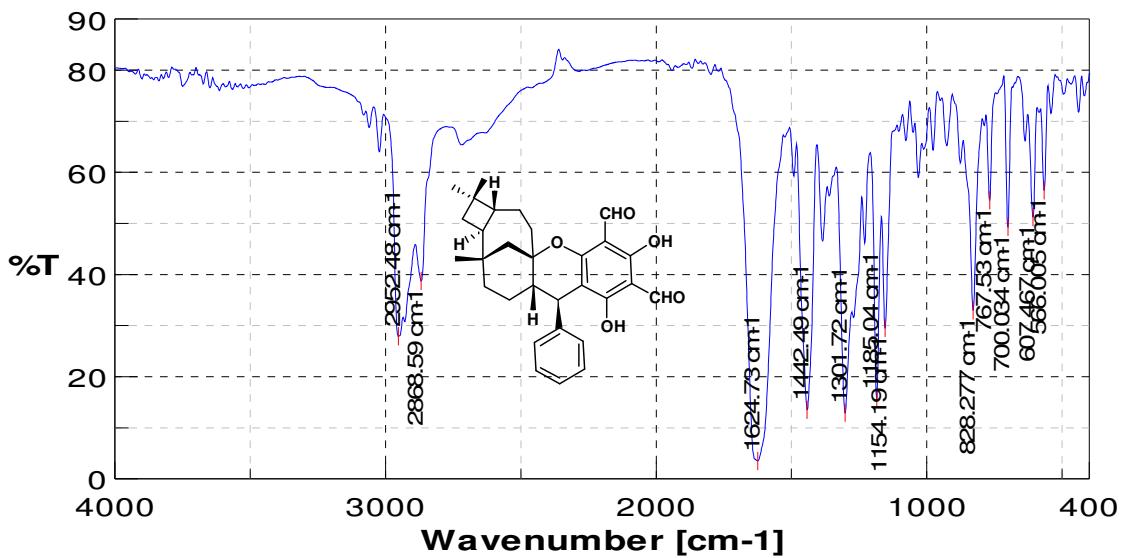
ROESY spectrum of **1** in CDCl_3

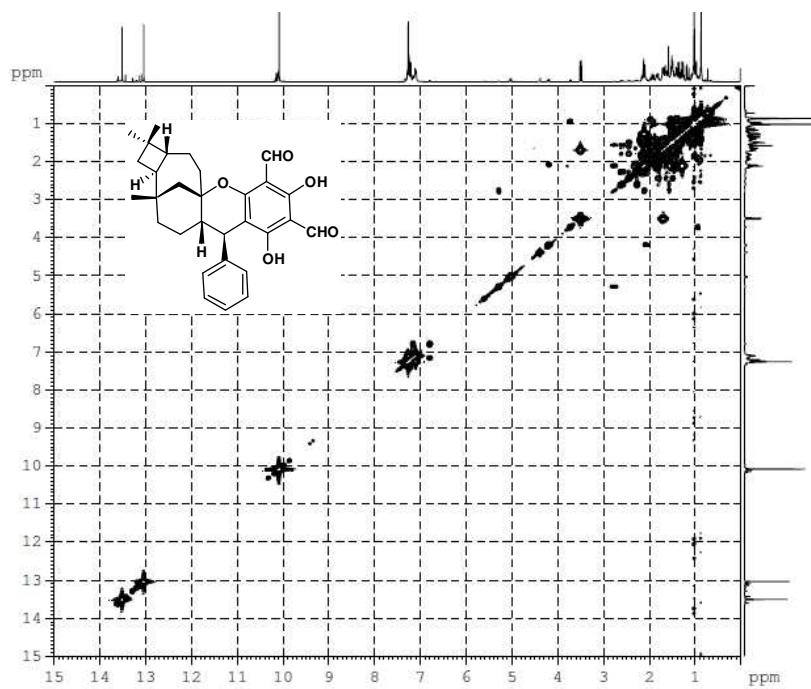
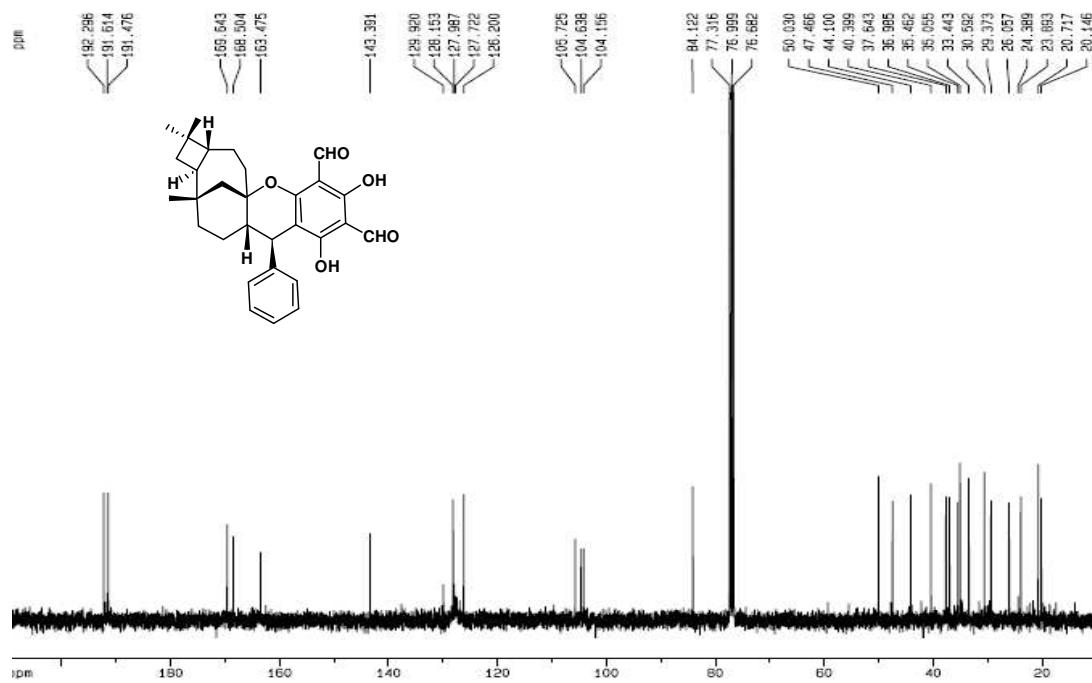


HR-EI-MS spectrum of **1**

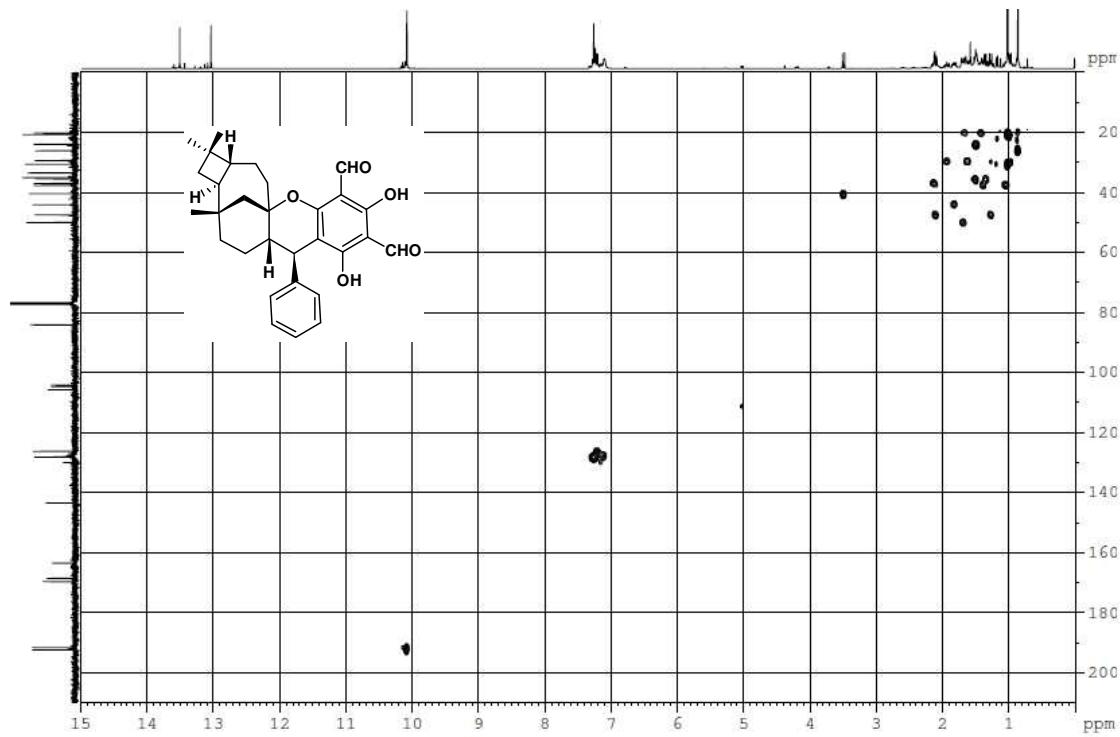


UV spectrum of **2** (CHCl_3)

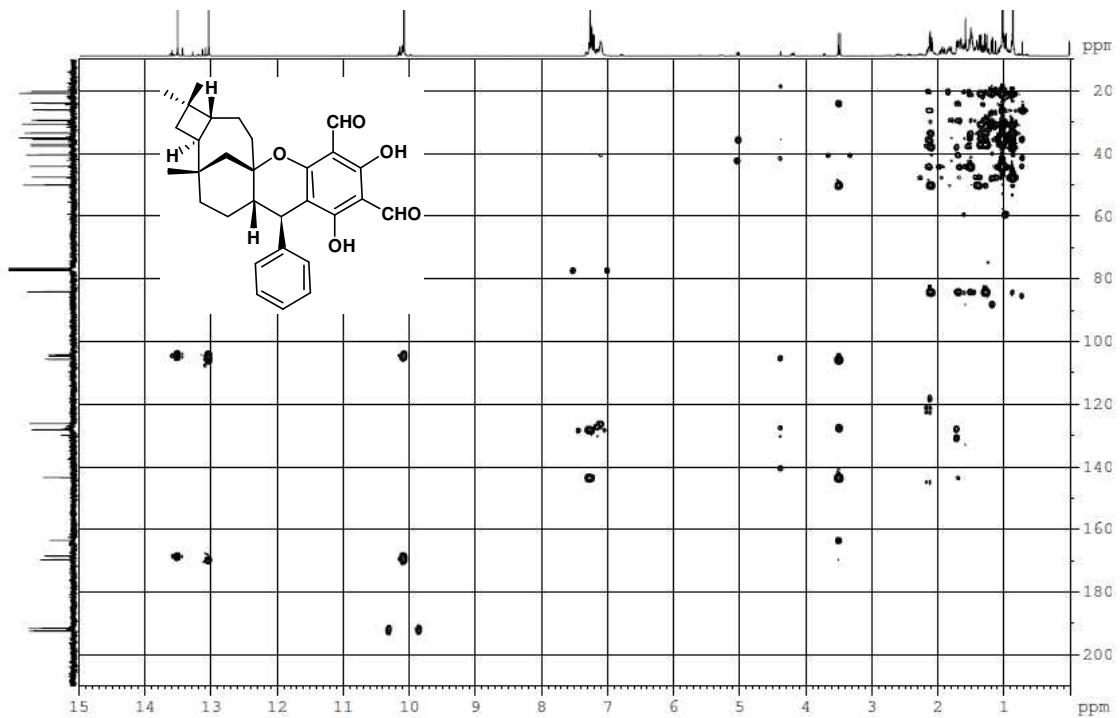




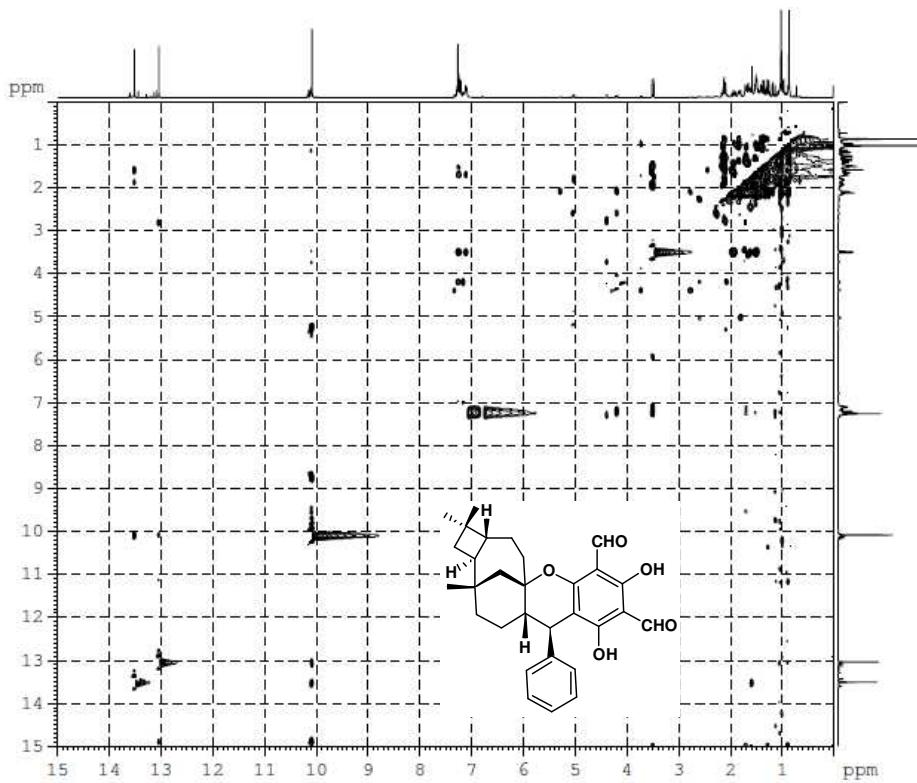
¹H-¹H COSY spectrum of **2** in CDCl₃



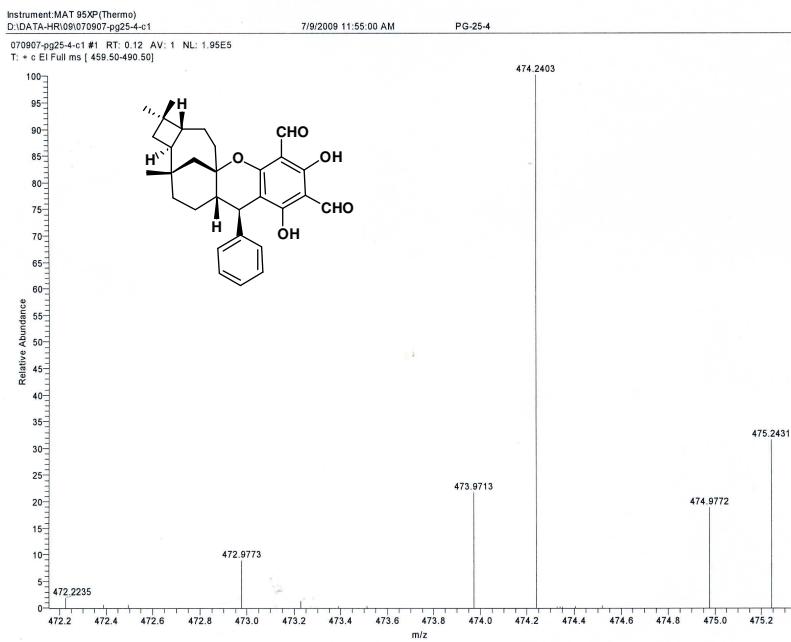
HSQC spectrum of **2** in CDCl_3



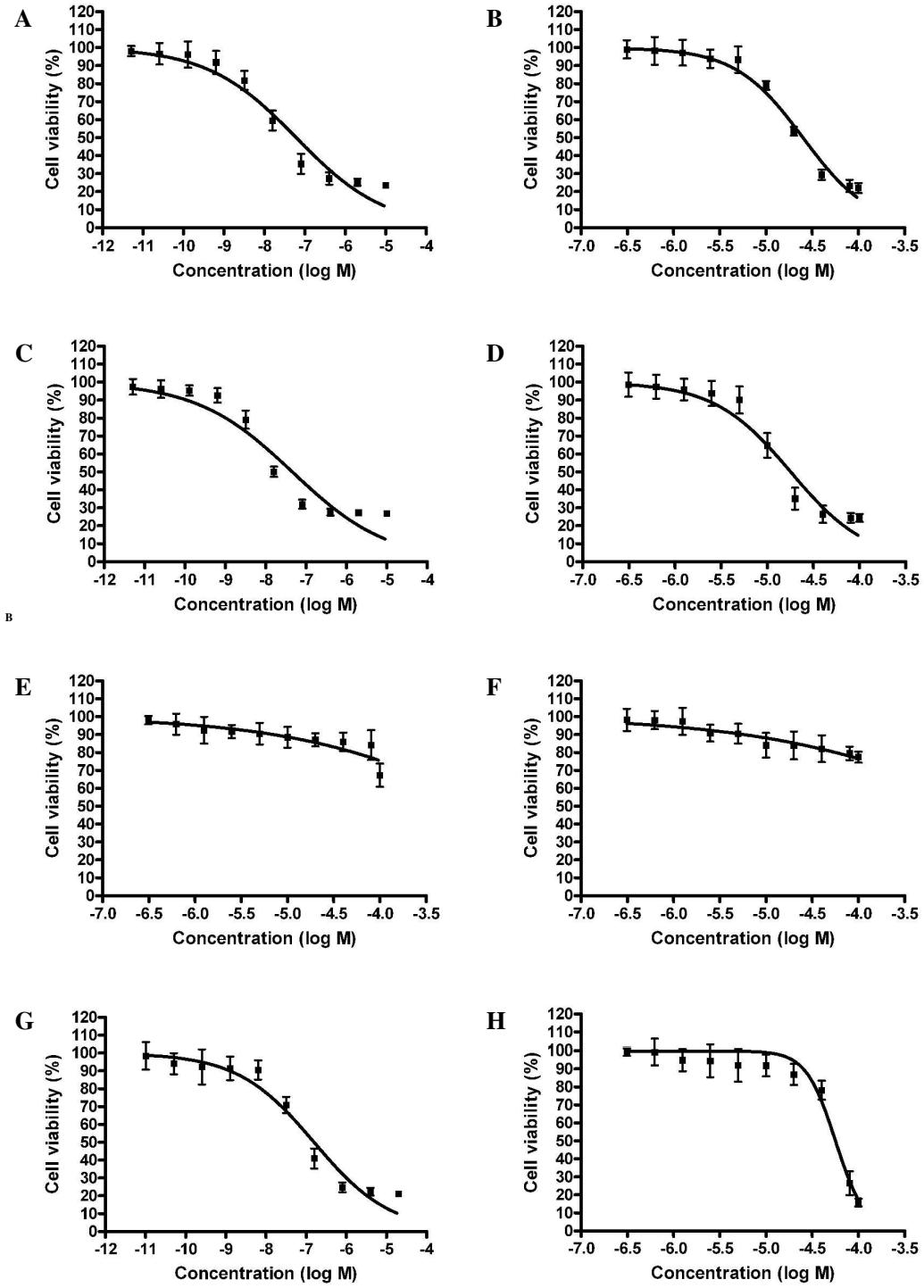
HMBC spectrum of **2** in CDCl_3



ROESY spectrum of **2** in CDCl_3



HR-EI-MS spectrum of **2**



Cell survival was determined by MTT assay. The survival curves of HepG2 cells at the different concentrations of **1** (A), **2** (C), **3** (E) and **4** (G). The survival curves of HepG2/ADM cells at the different concentration of **1** (B), **2** (D), **3** (F) and **4** (H). Data points are the means \pm SD of six replicates. Representative curves of three independent experiments are shown.