



Title	Isolation of rugosin A, B and related compounds as dipeptidyl peptidase-IV inhibitors from rose bud extract powder
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Supplementary material

Isolation of rugosin A, B and related compounds as dipeptidyl peptidase-IV inhibitors from rose bud extract powder

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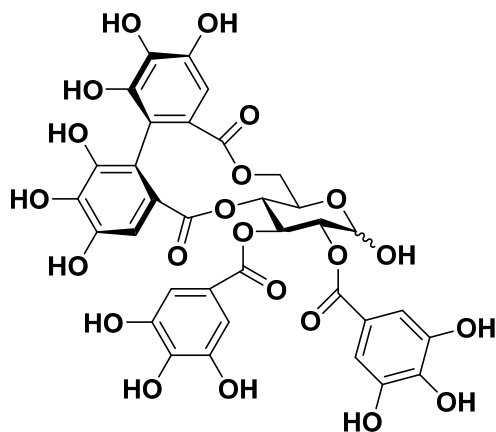
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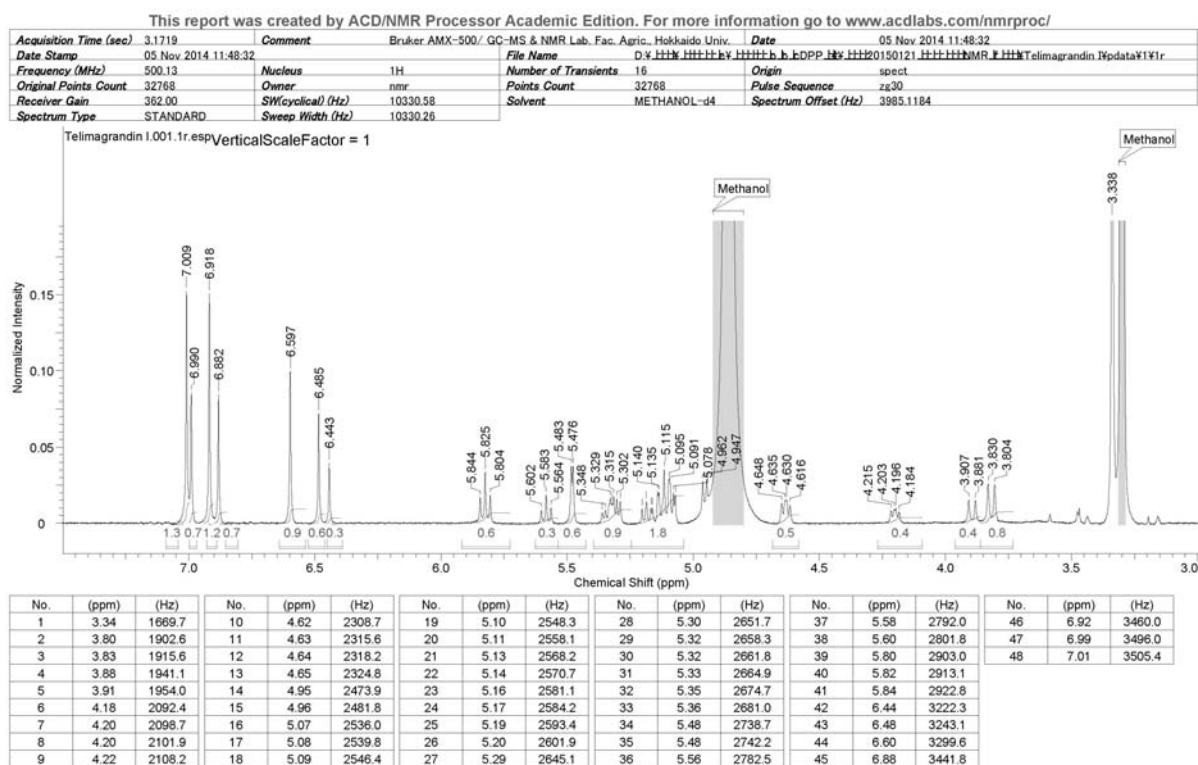
junk@chem.agr.hokudai.ac.jp (J. Kawabata)

¹ H NMR spectrum data of the isolated compounds in CD ₃ OD	2
Tellimagrandin I (1 , mixture of α, β-anomer)	2
Rugosin B (2)	3
Heterophyllin A (3)	4
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Tellimagrandin I (**1**, mixture of α , β -anomer)

Tellimagrandin I (1)

¹H NMR (500 MHz, CD₃OD, rt) α-anomer: 3.82 (1H, d, *J* = 12.8 Hz), 4.63 (1H, dd, *J* = 6.9, 9.9 Hz), 5.09 (1H, dd, *J* = 3.7, 9.9 Hz), 5.12 (1H, dd, *J* = 9.9, 10.0 Hz), 5.31 (1H, dd, *J* = 6.9, 12.8 Hz), 5.48 (1H, d, *J* = 3.7 Hz), 5.83 (1H, dd, *J* = 9.9, 10.0 Hz), 6.49 (1H, s), 6.60 (1H, s), 6.92 (2H, s), 7.01 (2H, s) ppm; β-anomer: 3.89 (1H, d, *J* = 12.9 Hz), 4.20 (1H, dd, *J* = 6.3, 9.5 Hz), 4.95 (1H, d, *J* = 7.9 Hz), 5.14 (1H, t, *J* = 10.2 Hz), 5.19 (1H, t, *J* = 8.9 Hz), 5.34 (1H, dd, *J* = 6.3, 13.2 Hz), 5.58 (1H, t, *J* = 9.6 Hz), 6.44 (1H, s), 6.60 (1H, s), 6.88 (2H, s), 6.99 (2H, s) ppm



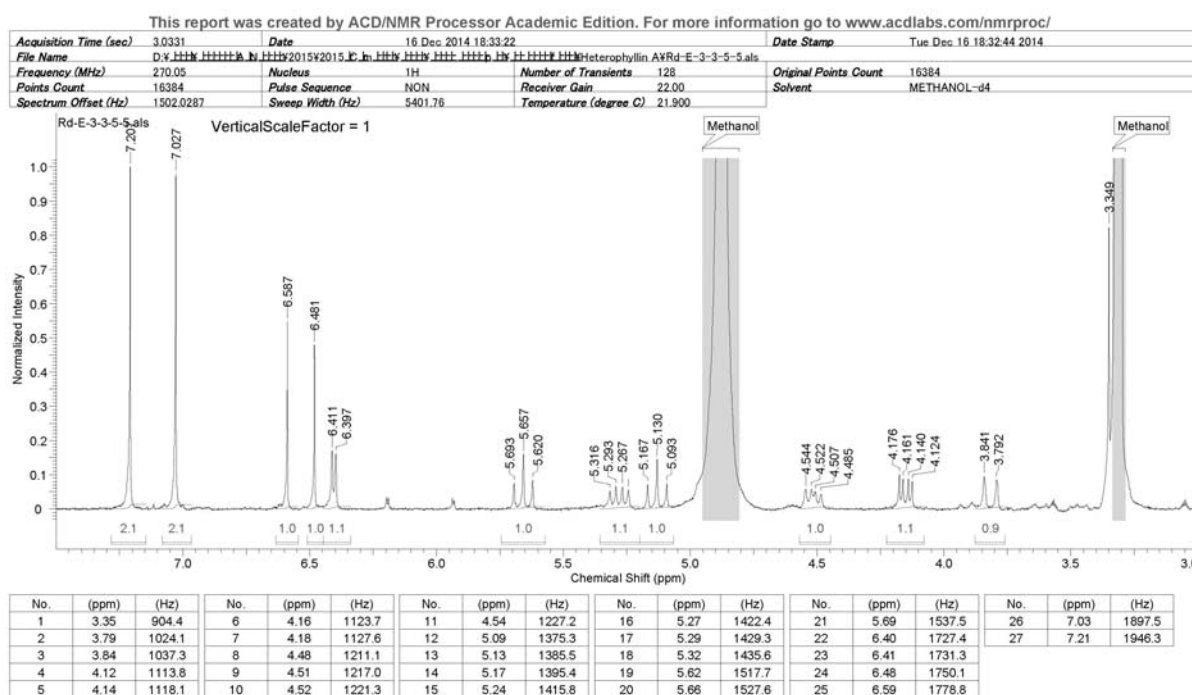
Rugosin B (2)

The chemical structure of Rugosin B (2) is a complex polycyclic molecule. It features a central core with multiple hydroxyl groups and a carboxylic acid group. The structure is highly branched and includes several fused and linked rings, including a benzene ring with multiple hydroxyl groups and a carboxylic acid group. The molecule is shown in a 2D representation with stereochemistry indicated by wedges and dashes.

β-anomer: 3.83 (1H, d, J = 13.2 Hz), 4.16 (1H, dd, J = 6.6, 10.1 Hz), 4.93 (1H, d, J = 8.2 Hz), 5.05-5.10 (1H, m), 5.16-5.27 (2H, m), 5.56 (1H, t, J = 9.8 Hz), 6.20 (1H, s), 6.46 (1H, s), 6.89 (2H, s), 7.00 (2H, s), 7.09 (1H, s) ppm



¹H-NMR (270 MHz, CD₃OD, rt): 3.82 (1H, d, J = 13.3 Hz), 4.15 (1H, dd, J = 4.1, 9.9 Hz), 4.52 (1H, dd, J = 6.2, 10.2 Hz), 5.13 (1H, dd, J = 10.0, 10.2 Hz), 5.28 (1H, dd, J = 6.2, 13.3 Hz), 5.66 (1H, dd, J = 9.9, 10.0 Hz), 6.40 (1H, d, J = 4.1 Hz), 6.48 (1H, s), 6.59 (1H, s), 7.03 (2H, s), 7.21 (2H, s)



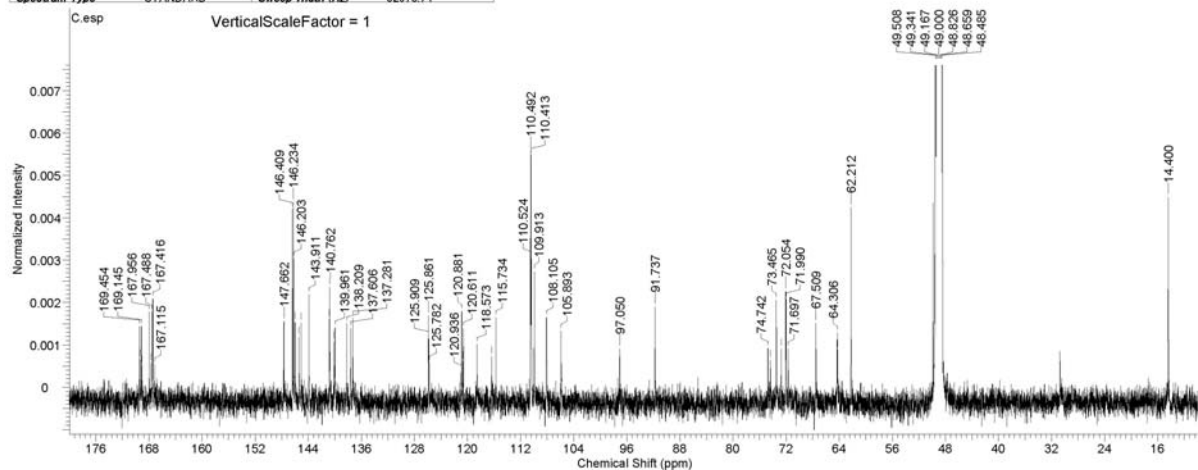
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This report was created by ACD/NMR Processor Academic Edition. For more information go to www.acdlabs.com/nmrproc/

Acquisition Time (sec)	3.1719	Comment	Bruker AMX-500/- GC-MS & NMR Lab. Fac. Agric. Hokkaido Univ.		Date	17 Feb 2015 10:01:52	
Date Stamp	17 Feb 2015 10:01:52	File Name	D:\114\114\data\114\1r				
Frequency (MHz)	500.13	Nucleus	1H	Number of Transients	16	Origin	spect
Original Points Count	32768	Owner	nmr	Points Count	32768	Pulse Sequence	zg30
Receiver Gain	256.00	SW(cyclical) (Hz)	10330.58	Solvent	METHANOL-d4	Spectrum Offset (Hz)	3990.4348
Spectrum Type	STANDARD	Sweep Width (Hz)	10330.26	Temperature (degree C)	24.200		

No.	(ppm)	(Hz)	No.	(ppm)	(Hz)	No.	(ppm)	(Hz)	No.	(ppm)	(Hz)	No.	(ppm)	(Hz)	No.	(ppm)	(Hz)
1	1.18	590.5	11	4.15	2073.2	21	4.59	2296.0	31	5.19	2594.6	41	5.54	2768.9	51	6.89	3447.1
2	1.20	597.7	12	4.15	2077.6	22	4.60	2303.0	32	5.20	2601.2	42	5.56	2778.7	52	6.93	3463.8
3	1.21	604.0	13	4.17	2084.5	23	4.94	2468.8	33	5.20	2603.1	43	5.58	2788.5	53	7.00	3498.8
4	1.21	604.7	14	4.18	2088.9	24	5.04	2518.6	34	5.21	2607.2	44	5.78	2889.1	54	7.01	3508.2
5	1.22	611.0	15	4.18	2091.4	25	5.06	2528.7	35	5.23	2614.1	45	5.80	2899.1	55	7.05	3524.9
6	3.74	1869.5	16	4.19	2095.9	26	5.08	2538.8	36	5.24	2618.2	46	5.82	2908.9			
7	3.76	1881.5	17	4.20	2098.4	27	5.08	2542.3	37	5.25	2625.2	47	6.16	3079.8			
8	3.81	1905.4	18	4.21	2103.1	28	5.10	2548.9	38	5.26	2632.1	48	6.17	3083.6			
9	3.84	1918.4	19	4.57	2286.3	29	5.10	2552.4	39	5.45	2725.4	49	6.46	3230.2			
10	4.14	2072.5	20	4.59	2293.2	30	5.17	2585.8	40	5.46	2729.2	50	6.50	3251.0			

Acquisition Time (sec)	1.0027	Comment	Broker AMX-500/ GC-MS & NMR Lab. Fac. Agric. Hokkaido Univ.	Date	17 Feb 2015 15:26:08	
Date Stamp	17 Feb 2015 15:26:08		File Name	D:\H1\H111111\H111111.b.DDPP.b.NMR\RGugin B.ethyl ester\Cydata\111r		
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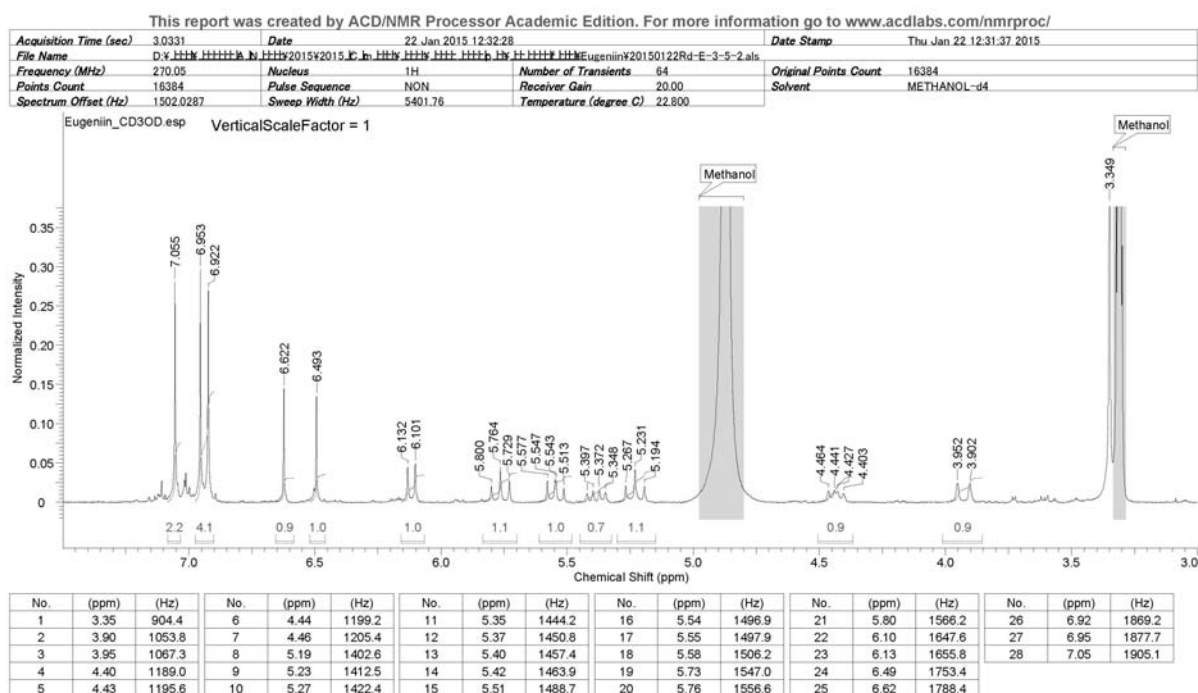


No.	(ppm)	(Hz)	No.	(ppm)	(Hz)	No.	(ppm)	(Hz)	No.	(ppm)	(Hz)	No.	(ppm)	(Hz)	No.	(ppm)	(Hz)
1	14.40	1810.9	12	64.31	8086.9	23	105.89	13316.8	34	120.67	15174.8	45	140.76	17702.0	56	147.71	18575.6
2	14.44	1815.9	13	67.51	8489.8	24	108.11	13595.1	35	120.88	15201.7	46	140.83	17709.9	57	167.12	21016.0
3	48.48	6097.3	14	71.70	9016.4	25	109.91	13822.4	36	120.94	15208.7	47	143.91	18097.9	58	167.42	21053.9
4	48.66	6119.2	15	71.99	9053.3	26	110.37	13879.3	37	125.78	15818.1	48	145.07	18243.5	59	167.49	21062.9
5	48.83	6140.2	16	72.05	9061.3	27	110.41	13885.3	38	125.86	15828.0	49	145.40	18285.4	60	167.73	21092.8
6	49.00	6162.1	17	72.73	9146.1	28	110.49	13895.2	39	125.91	15834.0	50	145.99	18359.2	61	167.96	21121.7
7	49.17	6183.1	18	73.47	9238.8	29	110.52	13899.2	40	137.28	17264.1	51	146.20	18386.1	62	169.07	21261.4
8	49.34	6205.0	19	74.37	9352.5	30	115.73	14554.4	41	137.61	17305.0	52	146.23	18390.1	63	169.15	21271.3
9	49.51	6226.0	20	74.74	9399.4	31	116.33	14629.3	42	138.21	17380.8	53	146.39	18409.1	64	169.35	21297.3
10	62.21	7823.6	21	91.74	11536.6	32	118.57	14911.5	43	139.96	17601.2	54	146.41	18412.0	65	169.45	21310.2
11	64.22	8076.0	22	91.05	12204.8	33	120.61	15157.6	44	140.18	17628.2	55	147.66	18569.6			

The image displays a complex chemical structure of a polyphenolic compound, possibly a flavonoid glycoside. The central core consists of a benzopyrone system (chromone) with multiple hydroxyl groups. This core is linked via ester bonds to several gallic acid units (3,4,5-trihydroxybenzoic acid). The structure is highly symmetrical and complex, with multiple hydroxyl groups and ester linkages.

Eugeniin (5)

¹H-NMR (270 MHz, CD₃OD, rt): 3.93 (1H, d, J = 13.4 Hz), 4.43 (1H, dd, J = 6.6, 10.0 Hz), 5.23 (1H, dd, J = 9.5, 10.0 Hz), 5.38 (1H, dd, J = 6.6, 13.4 Hz), 5.55 (1H, dd, J = 8.2, 9.2 Hz), 5.76 (1H, dd, J = 9.2, 9.5 Hz), 6.11 (1H, d, J = 8.2 Hz), 6.49 (1H, s), 6.62 (1H, s), 6.92 (2H, s), 6.95 (2H, s), 7.06 (2H, s)



The chemical structure shows a central core consisting of a benzene ring fused to a five-membered lactone ring. This core is substituted with several hydroxyl groups and ester-linked side chains. One side chain is a large gallic acid moiety (3,4,5-trihydroxybenzoic acid) linked via an ester bond. Another side chain is a smaller catechol moiety (1,2-dihydroxybenzoic acid) also linked via an ester bond. The structure is highly complex and represents a typical polyphenolic compound found in plants.

¹H-NMR (500 MHz, CD₃OD, rt): 3.85 (1H, d, *J* = 13.3 Hz), 4.38 (1H, dd, *J* = 6.4, 9.9 Hz), 5.18 (1H, dd, *J* = 9.5, 9.9 Hz), 5.29 (1H, dd, *J* = 6.4, 13.3 Hz), 5.54 (1H, dd, *J* = 8.4, 9.4 Hz), 5.74 (1H, dd, *J* = 9.4, 9.5 Hz), 6.09 (1H, d, *J* = 8.4 Hz), 6.23 (1H, s), 6.49 (1H, s), 6.93 (2H, s), 6.95 (2H, s), 7.05 (2H, s), 7.07 (1H, s)



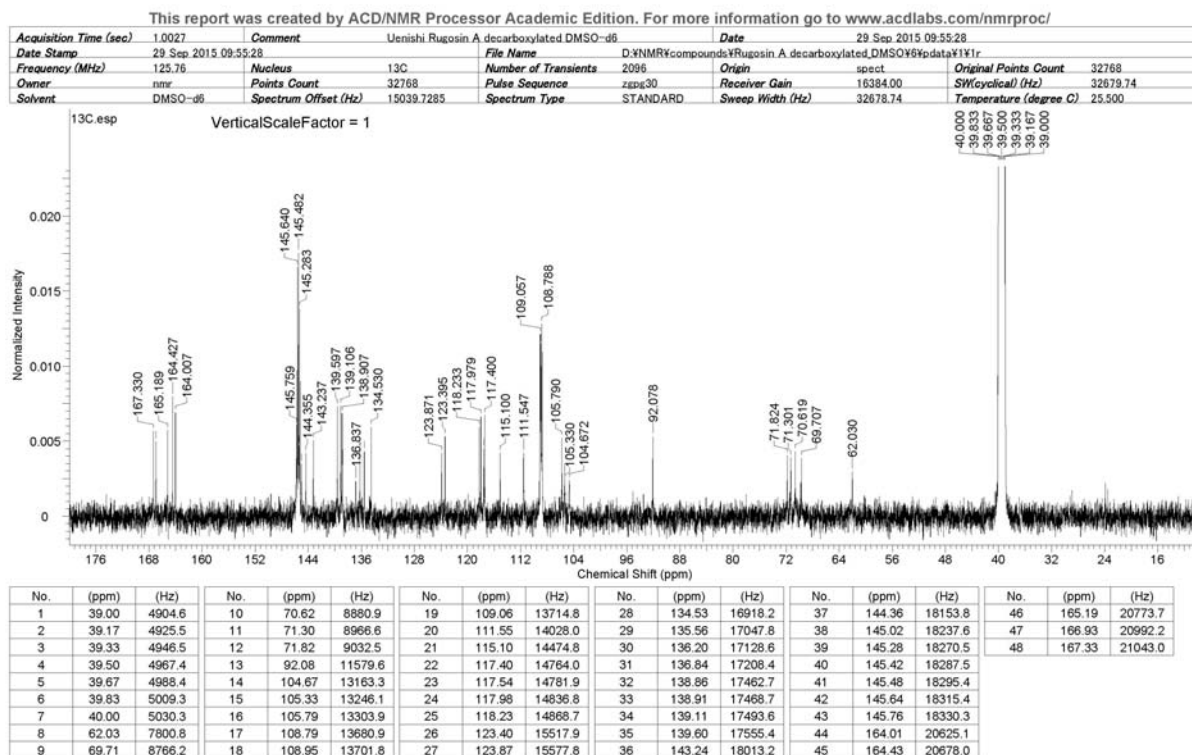
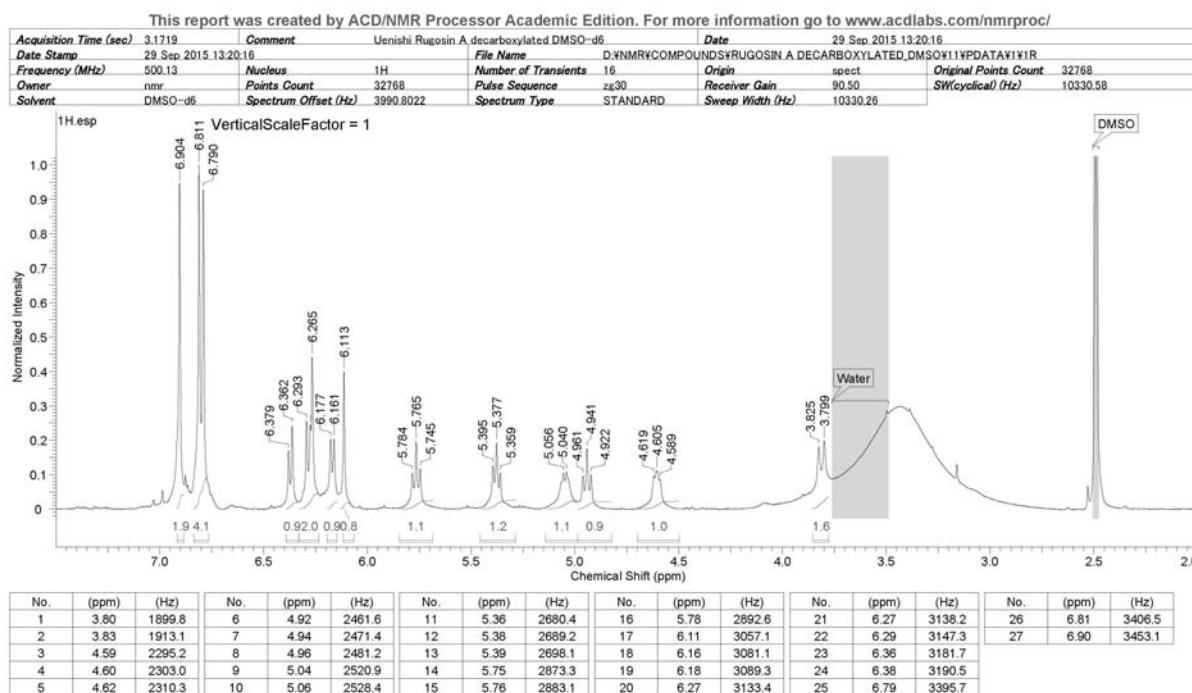
Oc1cc(O)c(O)c(O)c1Oc2cc(O)c(O)c(O)c2C(=O)O[C@H]3O[C@@H](OC(=O)c4cc(O)c(O)c(O)c4)[C@H](OC(=O)c5cc(O)c(O)c(O)c5)[C@H](OC(=O)c6cc(O)c(O)c(O)c6)[C@H](OC(=O)c7cc(O)c(O)c(O)c7)O3

¹H-NMR (500 MHz, CD₃OD, rt): 3.86 (1H, d, J = 13.4 Hz), 4.40 (1H, dd, J = 6.3, 10.0 Hz), 5.20 (1H, dd, J = 9.8, 10.0 Hz), 5.31 (1H, dd, J = 6.3, 13.4 Hz), 5.54 (1H, dd, J = 8.6, 9.3 Hz), 5.74 (1H, dd, J = 9.3, 9.8 Hz), 6.09 (1H, d, J = 8.6 Hz), 6.36 (1H, d, J = 9.2 Hz), 6.43 (1H, s), 6.44 (1H, d, J = 9.2 Hz), 6.50 (1H, s), 6.93 (2H, s), 6.95 (2H, s), 7.04 (2H, s)



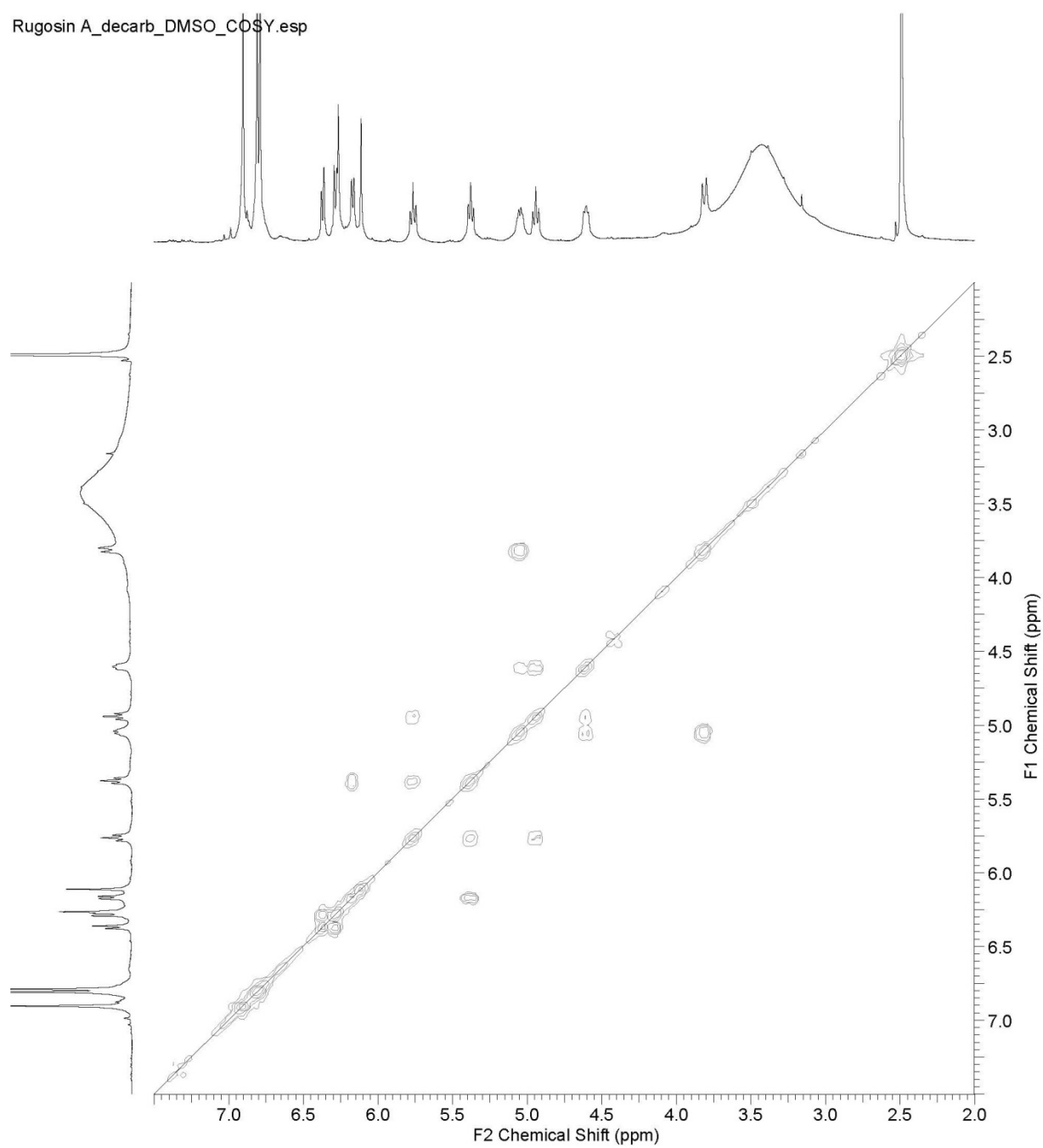
NMR spectra of rugosin A decarboxylated (7) in DMSO-*d*₆

¹H, ¹³C spectrum in DMSO-*d*₆



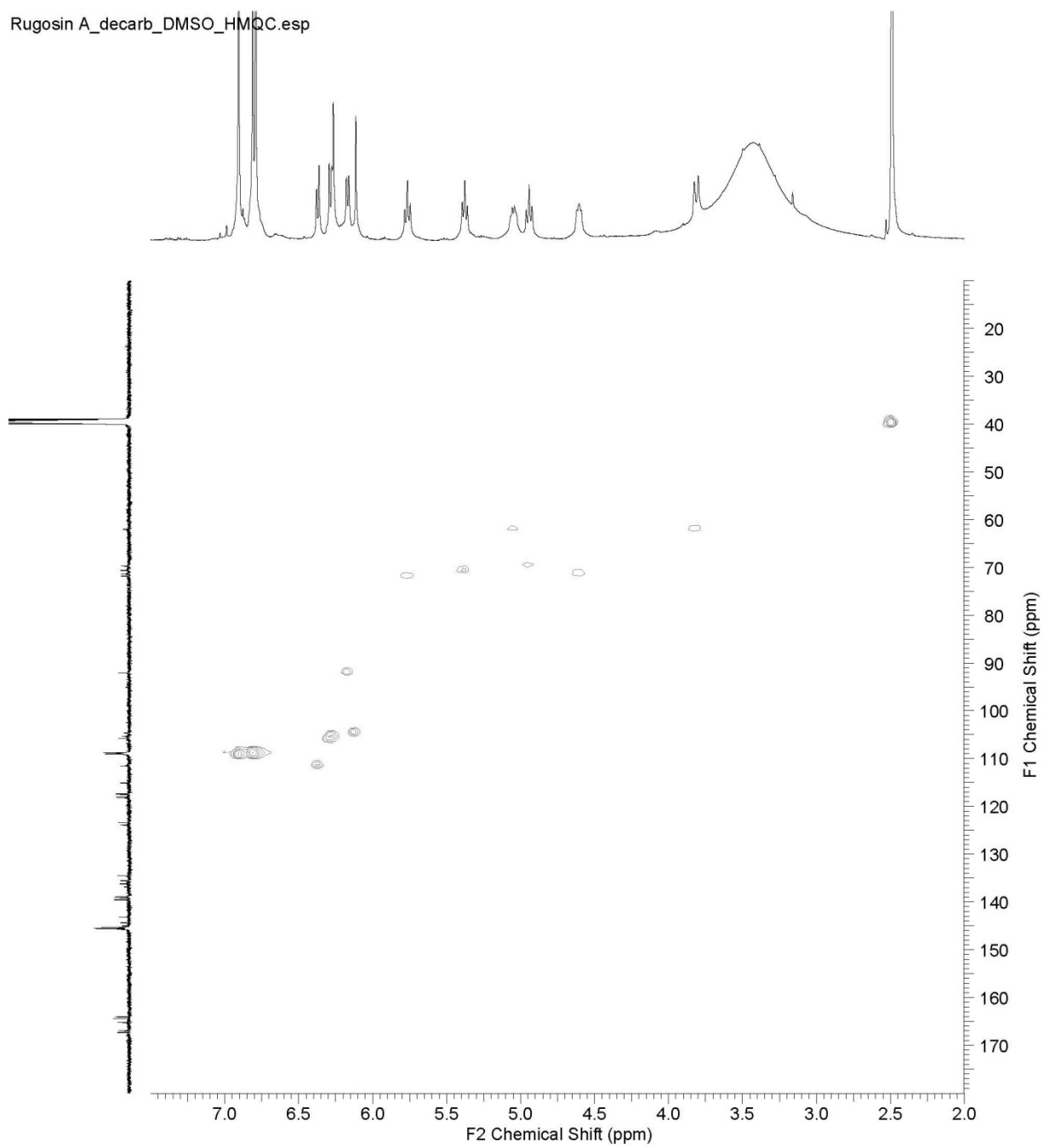
^1H - ^1H COSY spectrum in $\text{DMSO}-d_6$

Rugosin A_decarb_DMSO_COSY.esp



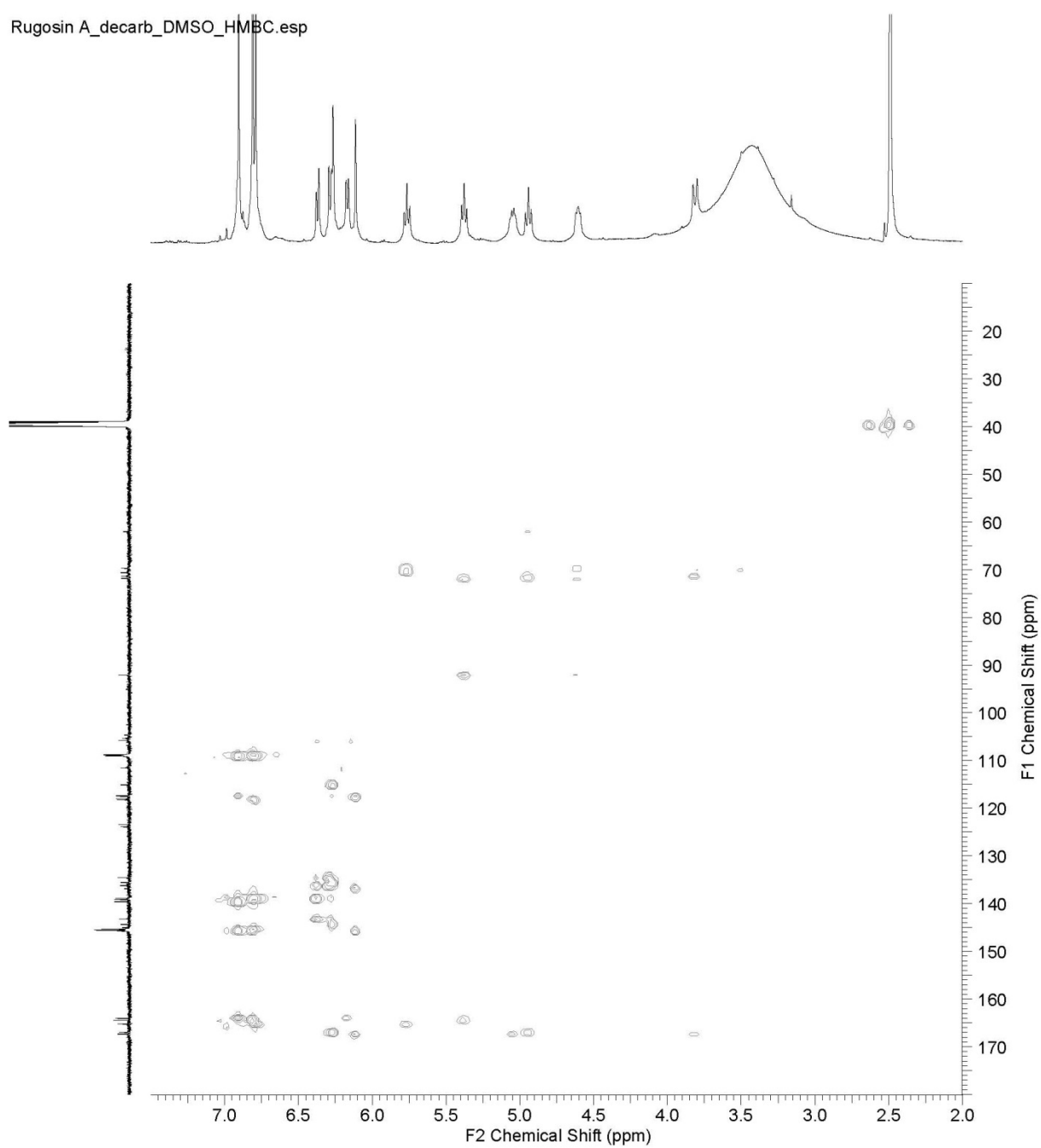
HMQC spectrum in DMSO-*d*₆

Rugosin A_decarb_DMSO_HMQC.esp



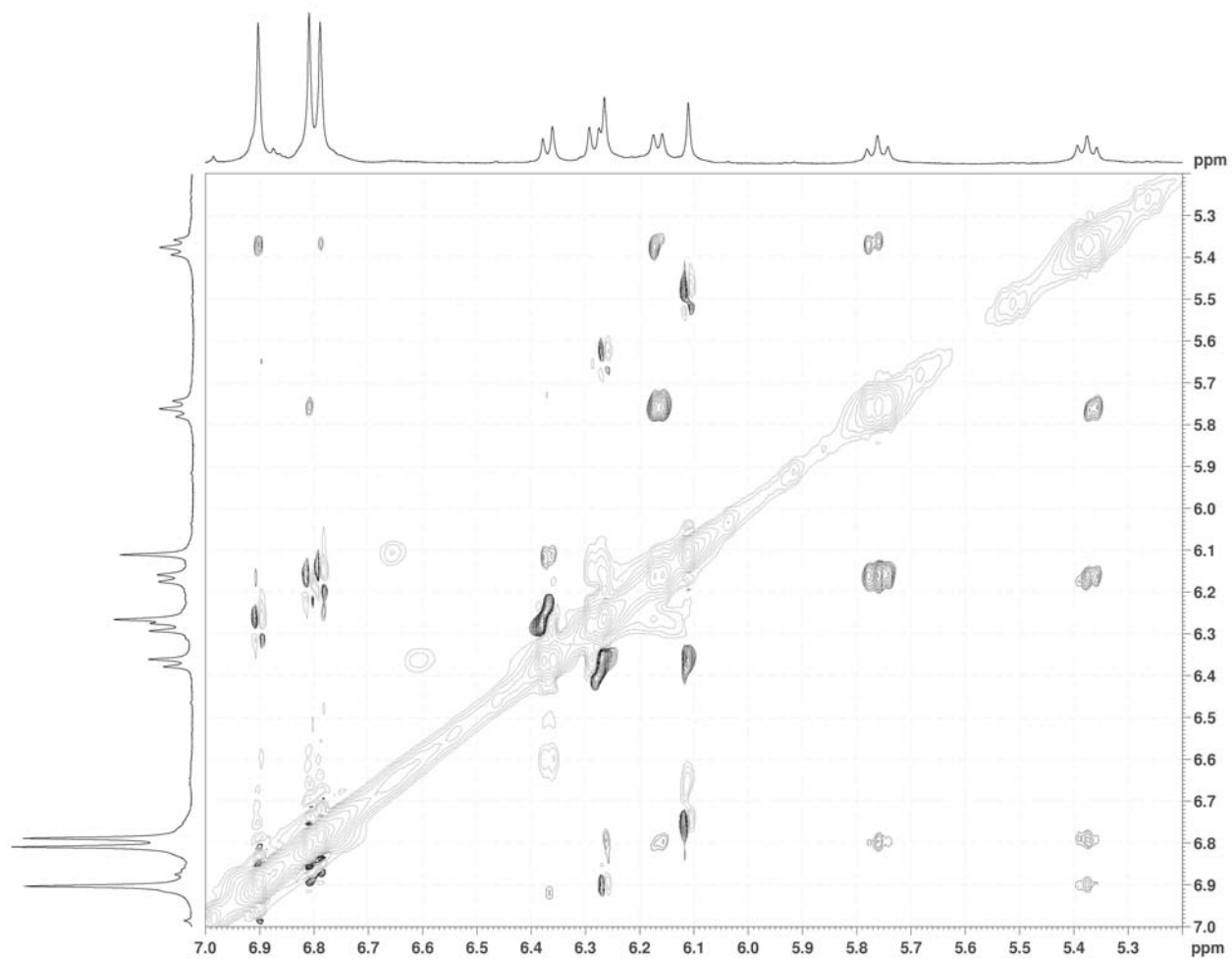
HMBC spectrum in DMSO- d_6

Rugosin A_decarb_DMSO_HMBC.esp



ROESY spectrum in DMSO- d_6

Uenishi
Rugosin A decarb. DMSO- d_6



UPLC-MS quantitation of rugosin A, B in rose buds extract powder

Condition

Column: Inertsustain C18 (ϕ 2.1×100 mm, GL Science Co.)

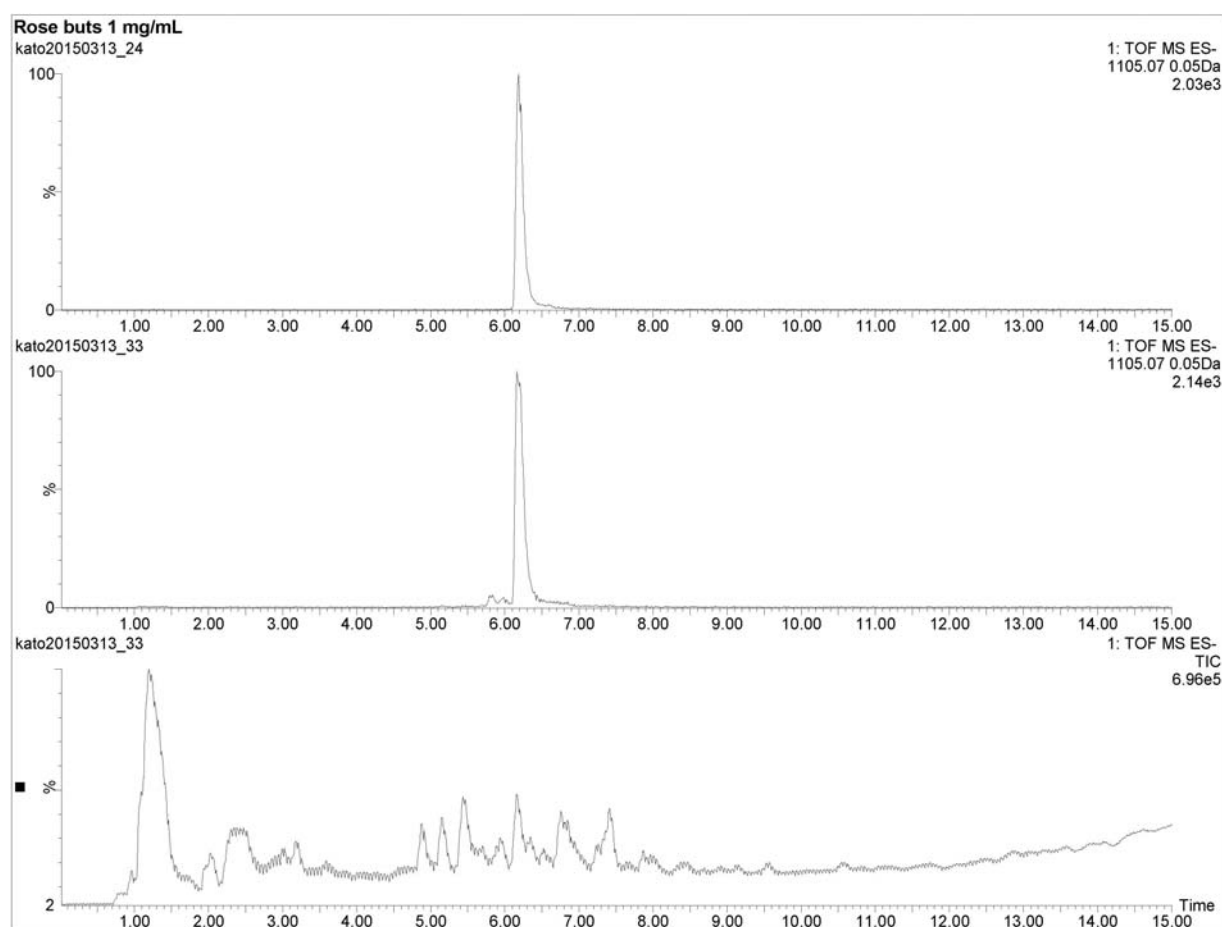
Mobile phase: gradient elution, 5% aq. MeCN with 0.1% TFA to 95% MeCN with 0.1% TFA (20 min)

Flow rate: 0.2 mL/min

Temperature: room temperature

Detection: negative mode ($[M-H]^-$)

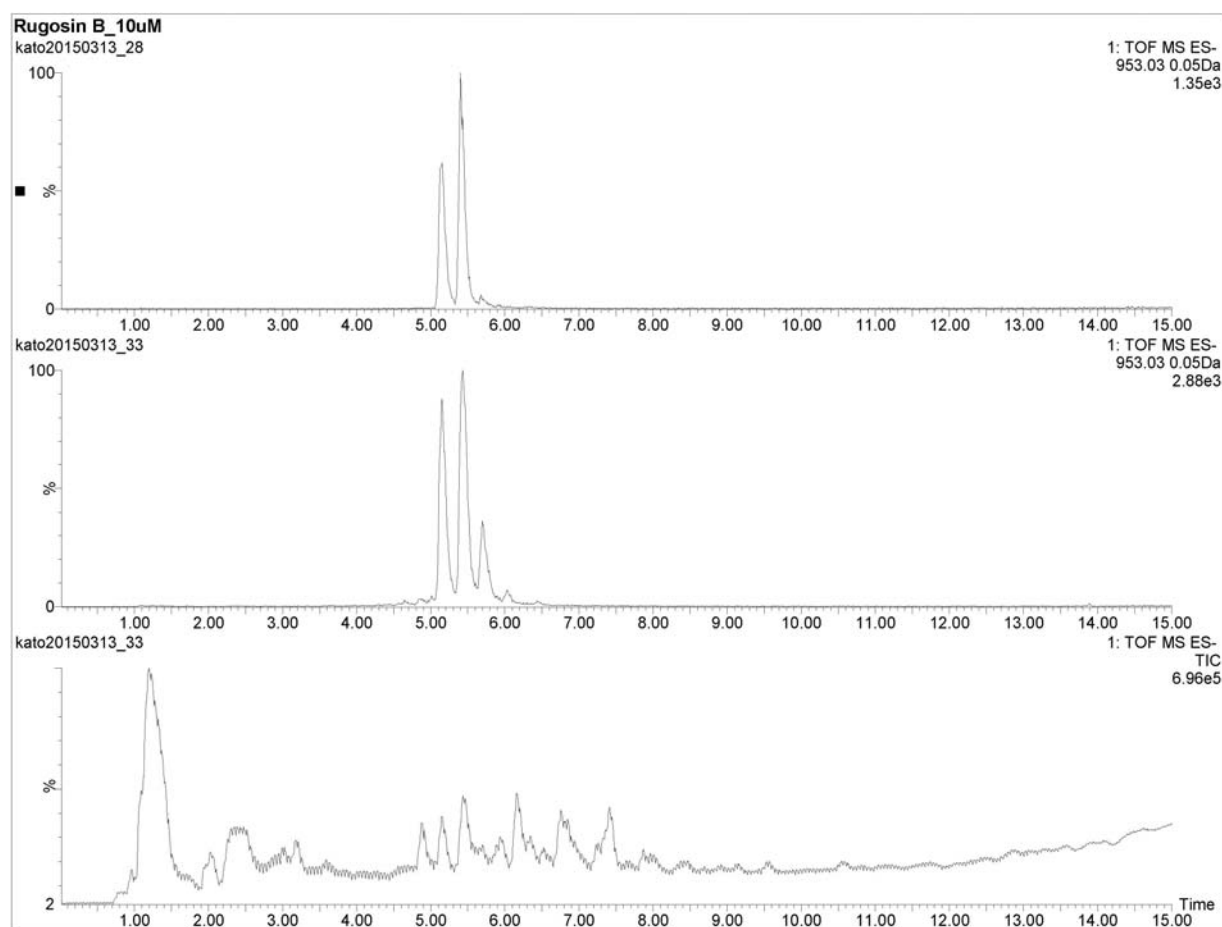
Samples: 4 μ L injection (Rose buds extract powder dissolved in water : 1 mg/mL)



Supplementary Figure 1. UPLC-MS analysis of rugosin A in the rose buds extract powder.

Top: Rugosin A (isolated sample); Middle: Selected ion chromatogram of rose buds extract powder;

Bottom: Total ion chromatogram of rose buds extract powder.



Supplementary Figure 2. UPLC-MS analysis of rugosin B in the rose butts extract powder.

Top: Rugosin B (isolated sample); Middle: Selected ion chromatogram of rose butts extract powder;

Bottom: Total ion chromatogram of rose butts extract powder.