

Supplementary Material

Identification of salicylates in willow bark (*Salix cortex*) for targeting peripheral inflammation

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Table of contents

1. High-Performance Liquid Chromatography (HPLC)	3
2. Isolation and identification of compounds 1-3 and 7 from SPE fraction F5	3
3. Isolation and identification of compounds 4 and 5 from SPE fraction F6	6
4. Isolation and identification of compound 6 from SPE fraction F7	8
5. Isolation and identification of compound 7 from subfraction F5-2.....	9
6. QTRAP-LC-MS/MS conditions for sugar determination	10
7. Absolute configuration of salicylates	11
8. Purification of fraction F7-4	11
9. NMR spectroscopic data	13

1. High-Performance Liquid Chromatography (HPLC)

For gradient and method development by means of analytical HPLC, the analytical 250 x 4.6 mm Luna® phenyl-hexyl column was used at room temperature. Therefore, SPE fractions F5, F6, and F7 were diluted in ACN/H₂O (v/v 70/30) to 1 mg/mL and injected into the HPLC system using the autosampler. The detection was performed by ELSD (evaporative light-scattering detector) and DAD (diode-array detector). Since all compounds were visible by UV, no ELSD was further used. Following settings were applied: 200 nm wavelength, 30 µL injection volume, and 1 mL/min flow rate. For chromatographic separation 0.1% formic acid in water (solvent A) and 0.1% formic acid in acetonitrile (solvent B) were used starting with a mixture of 5% B for 5 min, then increasing eluent B to 100% within 55 min, held isocratic for another 5 min, decreased again to 5% B in 4 min, and finally held at 5% B for 6 min.

Preparative HPLC separation of the three SPE fractions was performed on a preparative 250 x 21.2 mm Luna® phenyl-hexyl column using UV-detection at 200 nm, flow rate of 20 mL/min, and the same solvents as mentioned above. SPE fractions F5 and F7 were diluted in pure methanol to a final concentration of 10 mg/mL, whereas the concentration of F6 was 5 mg/mL in methanol. Semi-preparative HPLC fractionation was performed on a semi-preparative 250 x 10 mm, 5 µm, Luna® pentafluorophenyl column (Phenomenex Ltd., Aschaffenburg, Deutschland) for fractions F5-5, F5-2 and F7-8, and on a semi-preparative 250 x 10 mm, 5 µm, Luna® phenyl-hexyl column (Phenomenex Ltd., Aschaffenburg, Germany) for fractions F6-12 and F6-13 using 0.1% formic acid in water (solvent A) and 0.1% formic acid in methanol (solvent B). Following conditions were applied for the isolation and purification (purity ≥ 95% by NMR) of the compounds 1-7 by means of preparative (flow rate of 20 mL/min) and semi-preparative (flow rate of 4.7 mL/min) HPLC:

2. Isolation and identification of compounds 1-3 and 7 from SPE fraction F5

For the purification of the compounds 1-3 and 7, SPE fraction F5 was separated by means of preparative UV-HPLC ($\lambda = 200$ nm, injection volume: 500 µL, run time: 33 min). The chromatography started at 22% B, held for 3 min, increased in 10 min to 23.5% B, continued for 15 min at isocratic flow, and decreased to the initial conditions of 22% B, and finally held at 22% B for 3 min. Monitoring the UV signal at 200 nm, six fractions, F5-1 to F5-6, were collected, the solvent was removed, and used for further separation.

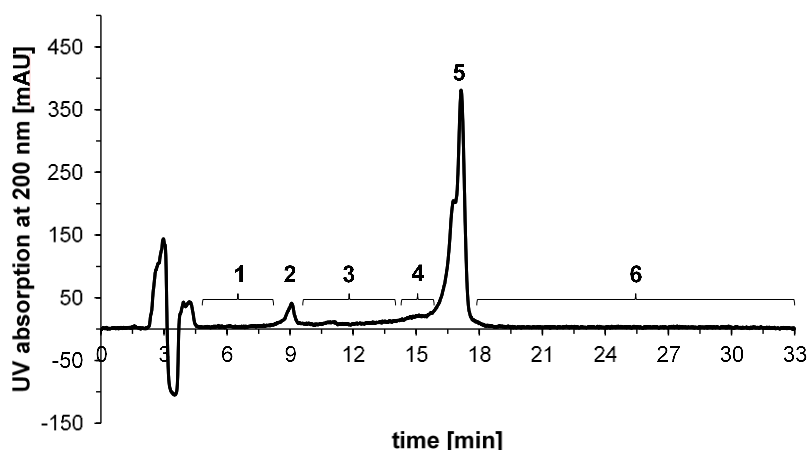


Figure S1: Preparative UV-HPLC chromatogram of SPE fraction F5 at 200 nm.

Further, fraction F5-5 (5 mg/mL in H₂O) was purified by means of semi-preparative UV-HPLC ($\lambda = 200$ nm, injection volume: 300 µL, run time: 65 min) and the following gradient: starting at 22% solvent B for 3 min, increased in 15 min to 27.7% B, held at 27.7% B for 2 min, increased to 52% B in 10 min, held at 52% B for 3 min, then increased further in 20 min to 57% B, held there for 2 min, and finally decreased in 3 min to 22% B, and held at 22% B for 2 min. 2'-O-acetylsalicortin (1), 3'-O-

acetylsalicortin (**2**) and 2'-*O*-acetylsalicin (**3**) were structurally elucidated in fractions F5-5-7 (**1**), F5-5-5 (**2**), and F5-5-3 (**3**) using LC-ToF-MS and 1D/2D-NMR experiments.

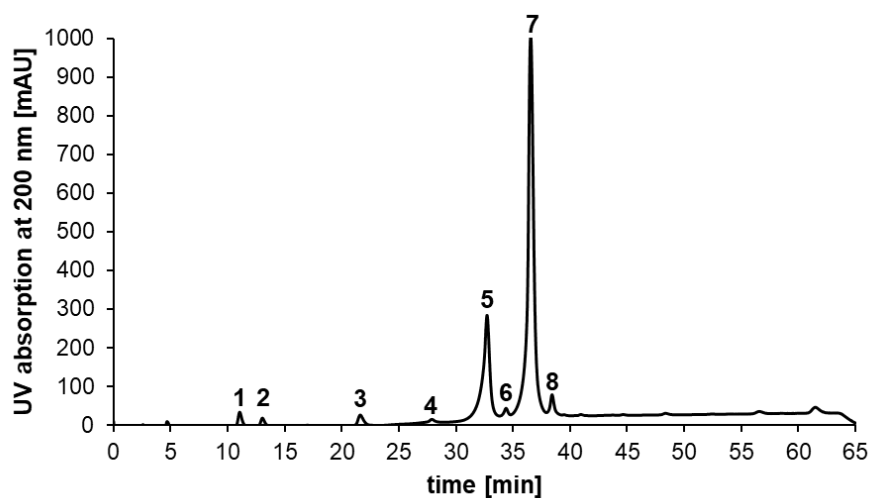


Figure S2: Semi-preparative UV-HPLC chromatogram of fraction F5-5 at 200 nm.

2'-*O*-acetylsalicortin (**1**)

$C_{22}H_{26}O_{11}$; UV (MeOH/H₂O, 1/1 v/v): λ_{\max} =204, 220, 272 nm; LC-TOF-MS (ESI): m/z 511.1455 [M+HCO₂H-H]⁻, 465.1434 [M-H]⁻; LC-MS/MS (DP = -160 V, CE = -80 V): m/z (%) 154.88 (100), 136.86 (78), 122.86 (57), 120.90 (22), 92.95 (22), 82.99 (19), 80.99 (10).

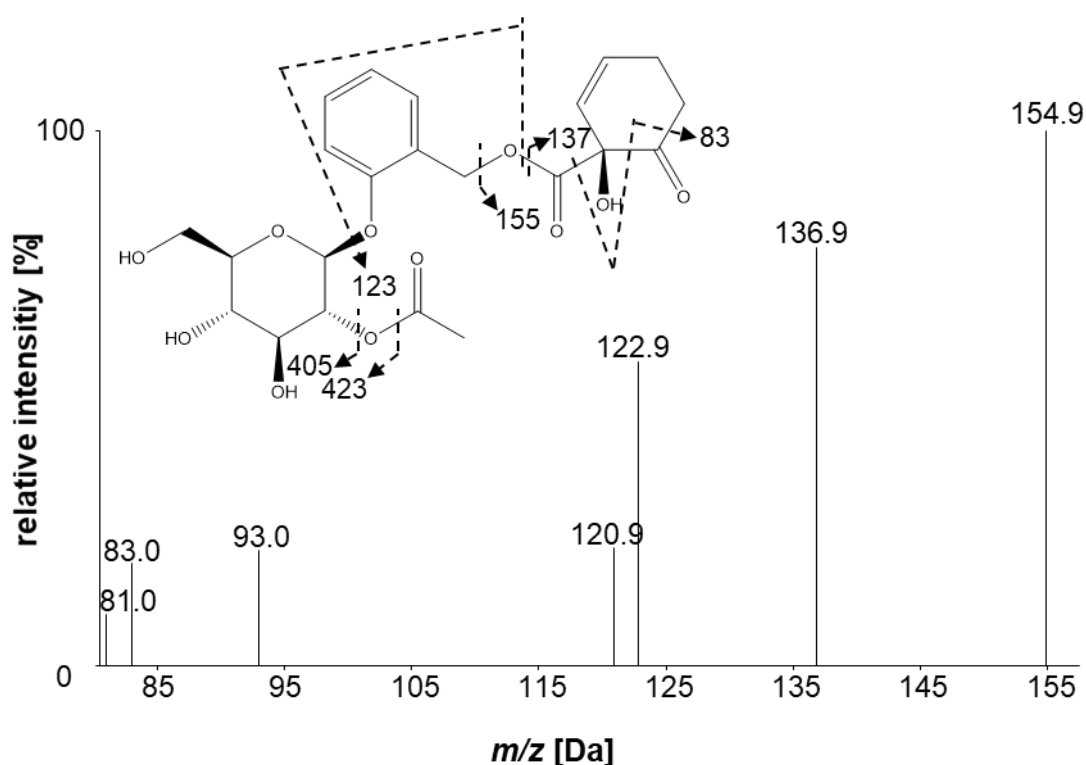


Figure S3: Centroided MS² spectrum of 464.9 Da showing the fragmentation pattern of 2'-*O*-acetylsalicortin (**1**).

3'-O-acetylsalicortin (2)

C₂₂H₂₆O₁₁; UV (MeOH/H₂O, 1/1 v/v): λ_{max} =200, 220, 272 nm; LC-TOF-MS (ESI): m/z 511.1458 [M+HCO₂H-H]⁻, 465.1421 [M-H]⁻; LC-MS/MS (DP = -80 V, CE = -66 V): m/z (%) 404.92 (41), 154.92 (100), 136.94 (76), 122.91 (61), 120.94 (19), 83.02 (19), 80.97 (5).

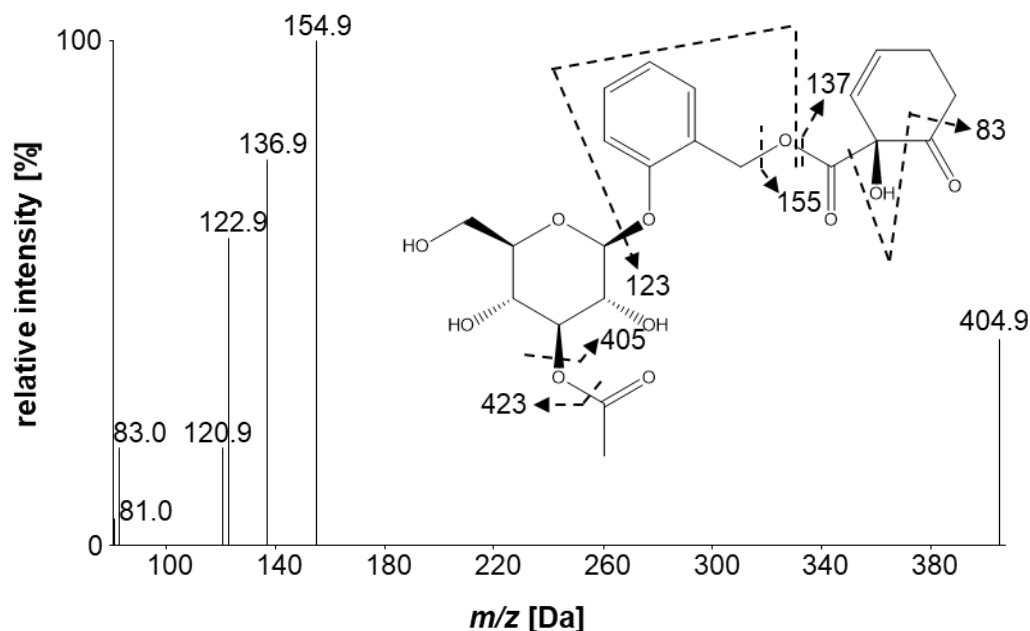


Figure S4: Centroided MS² spectrum of 465.0 Da showing the fragmentation pattern of 3'-O-acetylsalicortin (2).

2'-O-acetylsalicin (3)

C₁₅H₂₀O₈; UV (H₂O): λ_{max} = 204, 220, 268 nm; LC-TOF-MS (ESI): m/z 373.1130 [M+HCO₂H-H]⁻, 327.1070 [M-H]⁻; LC-MS/MS (DP = -5 V, CE = -76 V): m/z (%) 326.92 (100), 304.73 (73), 174.82 (32), 122.95 (87), 120.93 (8), 92.92 (3).

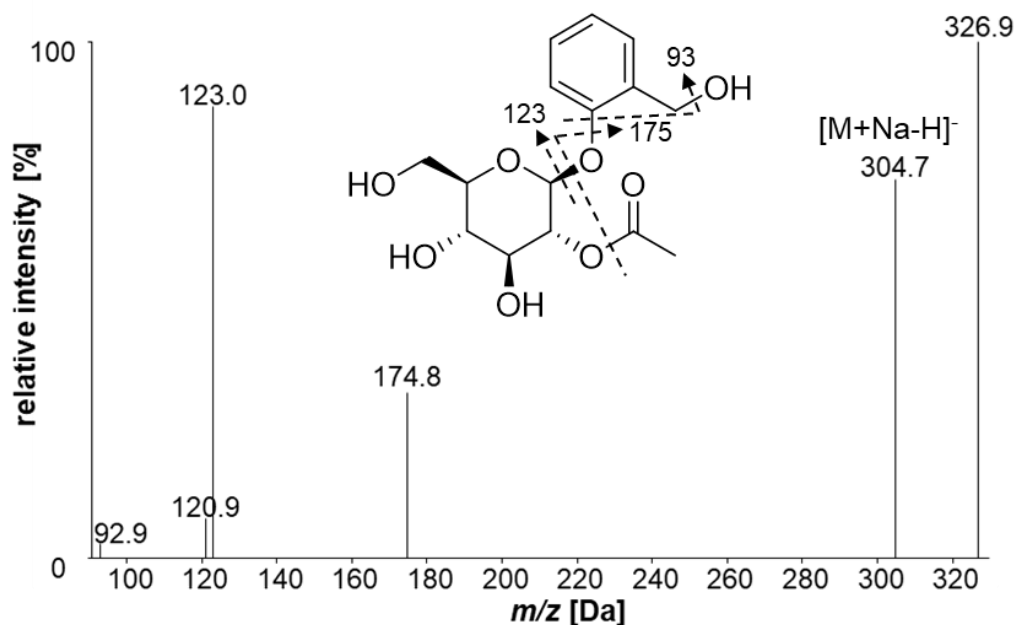


Figure S5: Centroided MS² spectrum of m/z 373.0 [M+HCO₂H-H]⁻ showing the fragmentation pattern of 2'-O-acetylsalicin (3).

3. Isolation and identification of compounds 4 and 5 from SPE fraction F6

SPE Fraction F6 was purified by means of preparative UV-HPLC ($\lambda = 200$ nm, injection volume: 600 μ L, run time: 35 min) and the following conditions: starting at 23% B, held isocratically for 3 min, increased in 21 min to 33% B, held at 33% B for 6 min, and decreased finally in 3 min to 23% B, which was held for 2 min. Fourteen fractions, F6-1 to F6-14, were collected, freed from the solvent, and used for further separation.

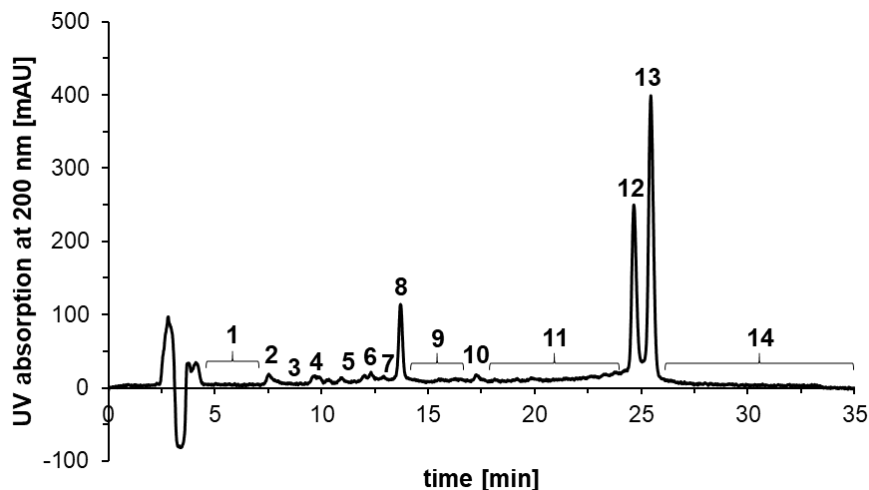


Figure S6: UV-HPLC chromatogram ($\lambda = 200$ nm) of SPE fraction F6 prepared from the methanol extract of *S. pentandra*.

Fraction F6-12 (10 mg/mL in MeOH/H₂O (v/v 1/1)) was purified by means of semi-preparative UV-HPLC ($\lambda = 200$ nm, injection volume: 150 μ L, run time: 50 min): the chromatography started at 20% B, held there for 3 min, increased in 43.5 min to 32% B, decreased again in 1.5 min to 20% B and held at 20% B for 2 min. 2',6'-O-diacetylsalicortin (**4**) was structurally elucidated in fraction 6-12-2 using LC-ToF-MS and 1D/2D-NMR experiments.

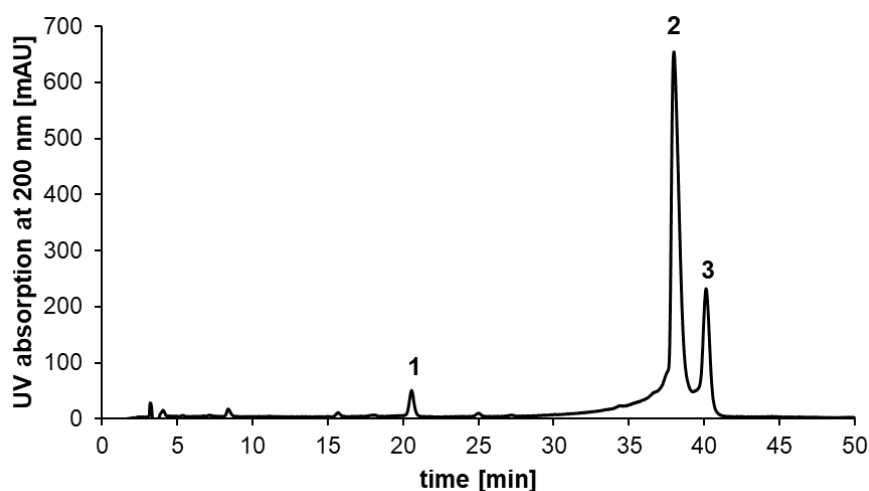


Figure S7: Semi-preparative UV-HPLC ($\lambda = 200$ nm) chromatogram of fraction F6-12.

2',6'-O-diacetylsalicortin (**4**)

C₂₄H₂₈O₁₂; UV (MeOH/H₂O, 1/1 v/v): λ_{max} =200, 272, 300 nm; LC-TOF-MS (ESI): m/z 553.1569 [M+HCO₂H-H]⁻, 507.1551 [M-H]⁻; LC-MS/MS (DP = -160 V, CE = -62 V): m/z (%) 155.00 (81), 136.90 (100), 122.89 (58), 120.91 (22), 92.94 (23), 82.94 (17), 80.98 (4).

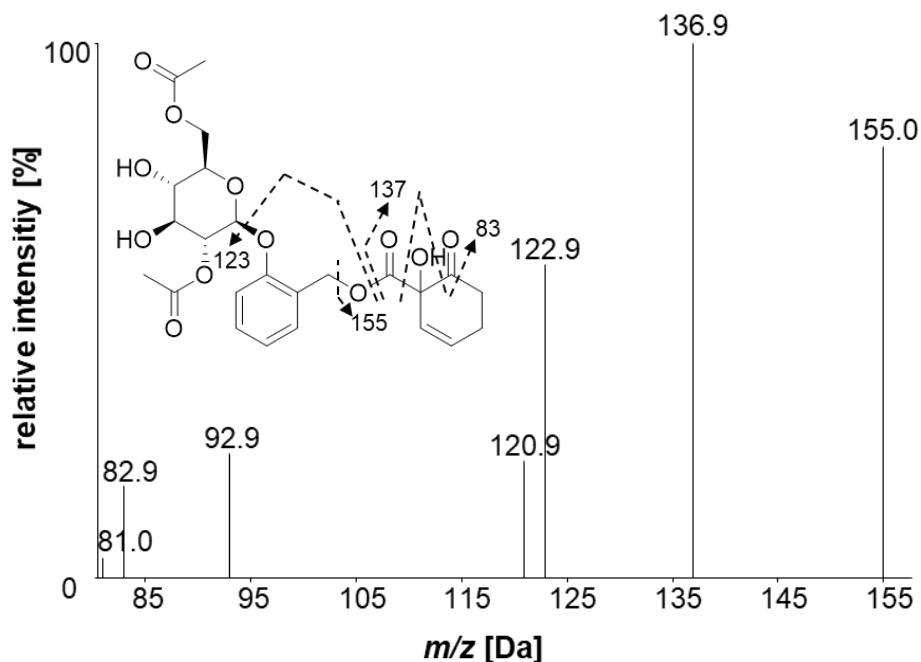


Figure S8: Centroided MS² spectrum of *m/z* 507.0 showing the fragmentation pattern of 2',6'-*O*-diacetylsalicortin (4).

Fraction F6-13 (10 mg/mL in MeOH/H₂O (v/v 1/1)) was purified by means of semi-preparative UV-HPLC ($\lambda = 200$ nm, injection volume: 200 μ L, run time: 34 min) and the following gradient: starting with an isocratic step at 27% B for 3 min, increased in 27 min to 32% B, finally decreased in 4 min to 27% B and held there. Lasiandrin (5) was structurally elucidated in fraction 6-13-2 using LC-ToF-MS and 1D/2D-NMR experiments.

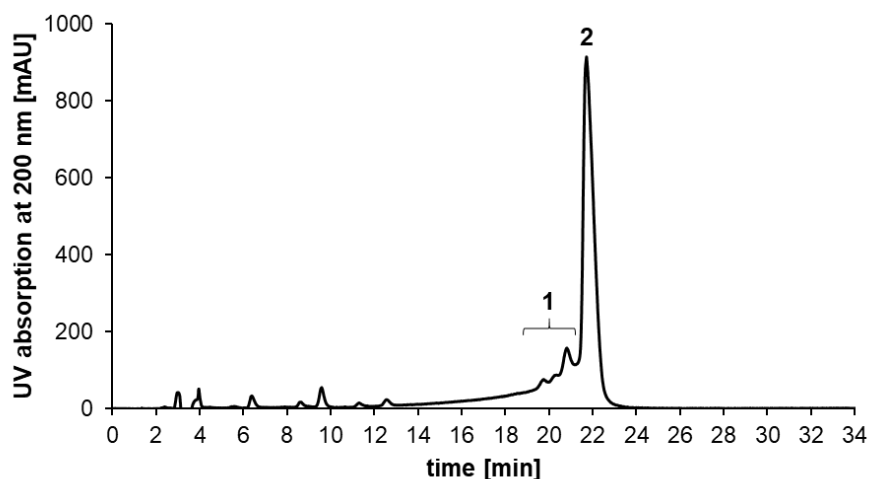


Figure S9: Semi-preparative UV-HPLC chromatogram of fraction F6-13 at 200 nm.

lasiandrin (5)

C₂₉H₃₂O₁₄; UV (MeOH/H₂O, 1/1 v/v): λ_{max} =200, 272, 300 nm; LC-TOF-MS (ESI⁻): *m/z* 649.1791 [M+HCO₂H-H]⁻, 603.1762 [M-H]⁻; LC-MS/MS (DP = -135 V, CE = -84 V): *m/z* (%) 464.99 (100), 154.90 (85), 136.91 (83), 110.95 (32), 108.93 (15), 92.95 (30), 82.98 (26), 80.96 (10).

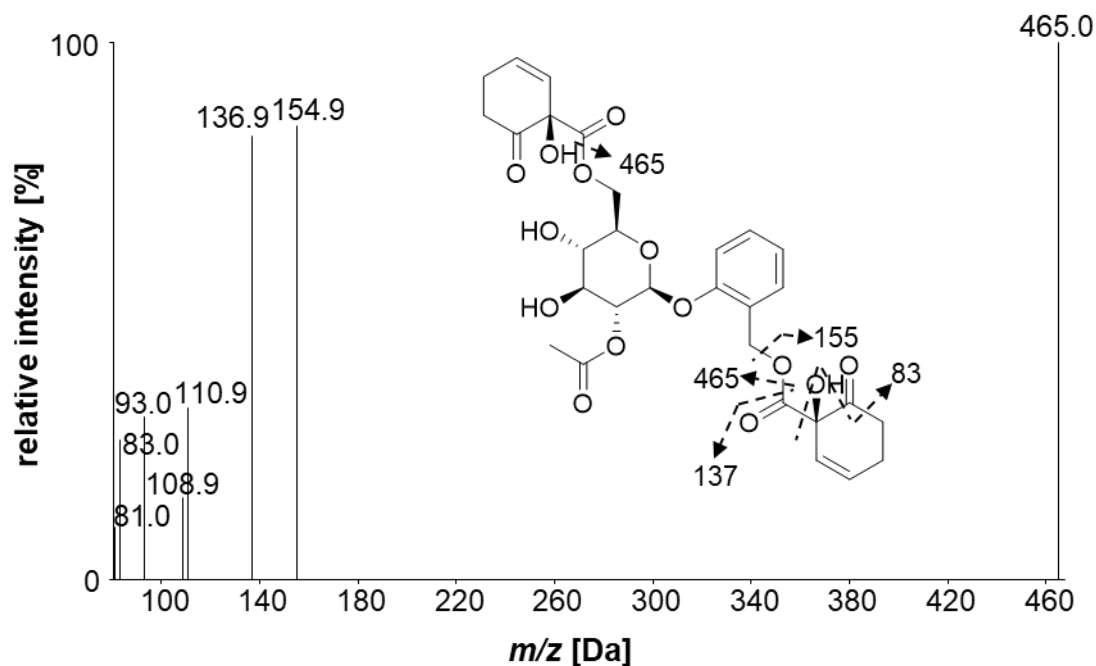


Figure S10: Centroided MS² spectrum of *m/z* 603.2 showing the fragmentation pattern of lasiandrin (5).

4. Isolation and identification of compound 6 from SPE fraction F7

SPE fraction F7 was purified by means of preparative UV-HPLC ($\lambda = 200$ nm, injection volume: 700 μ L, run time: 34 min): the chromatography started at 28% B, held there for 3 min, increased in 7 min to 32% B, isocratic for 2 min, increased in 14 min to 37% B, isocratic at 37% B for 2 min, decreased again in 3 min to 28% B, where it was held isocratic for 3 min. Fourteen fractions, F7-1 to F7-14, were collected, freed from the solvent, and used for further separation.

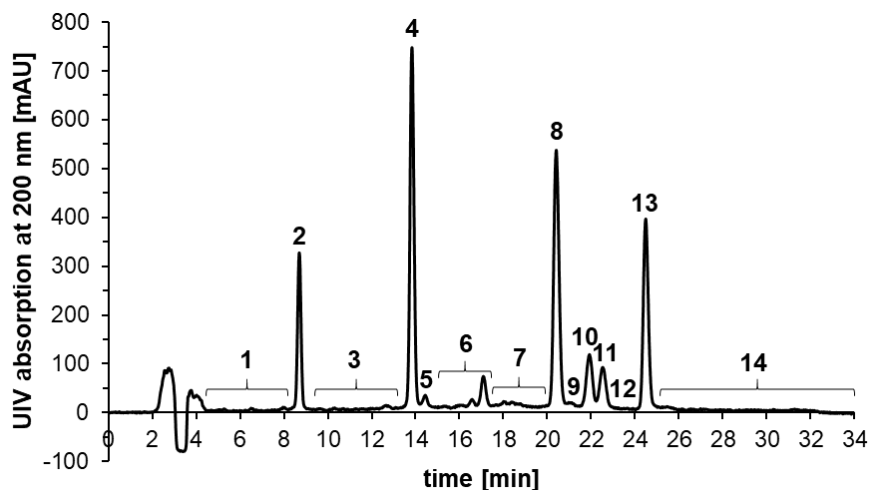


Figure S11: UV-HPLC chromatogram of SPE fraction F7 at 200 nm.

Fraction F7-8 was purified by means of semi-preparative UV-HPLC ($\lambda = 200$ nm, injection volume: 200 μ L, run time: 42 min) and the following chromatographic conditions: starting at 26% B, held there isocratic for 3 min, increased in 23 min to 62% B, held at 62% B for 10 min, decreased in 4 min to 26 % B, where it was held isocratic for 2 min. Tremulacin (6) was structurally characterized in fraction 7-8-4 using LC-ToF-MS and 1D/2D-NMR experiments.

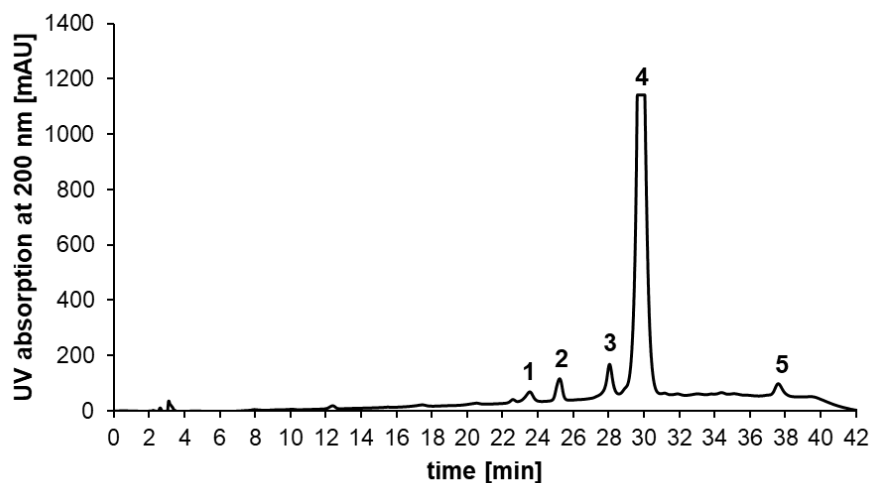


Figure S12: Semi-preparative UV-HPLC chromatogram of fraction F7-8 at 200 nm.

tremulacin (6)

$C_{27}H_{28}O_{11}$; UV (MeOH/H₂O, 1/1 v/v): λ_{max} =204, 236, 272 nm; LC-TOF-MS (ESI): m/z 573.1617 [M+HCO₂H-H]⁻, 527.1586 [M-H]⁻; LC-MS/MS (DP = -20 V, CE = -86 V): m/z (%) 404.95 (100), 154.90 (49), 136.90 (39), 122.92 (14), 120.94 (67), 82.97 (10), 80.97 (10), 76.97 (22).

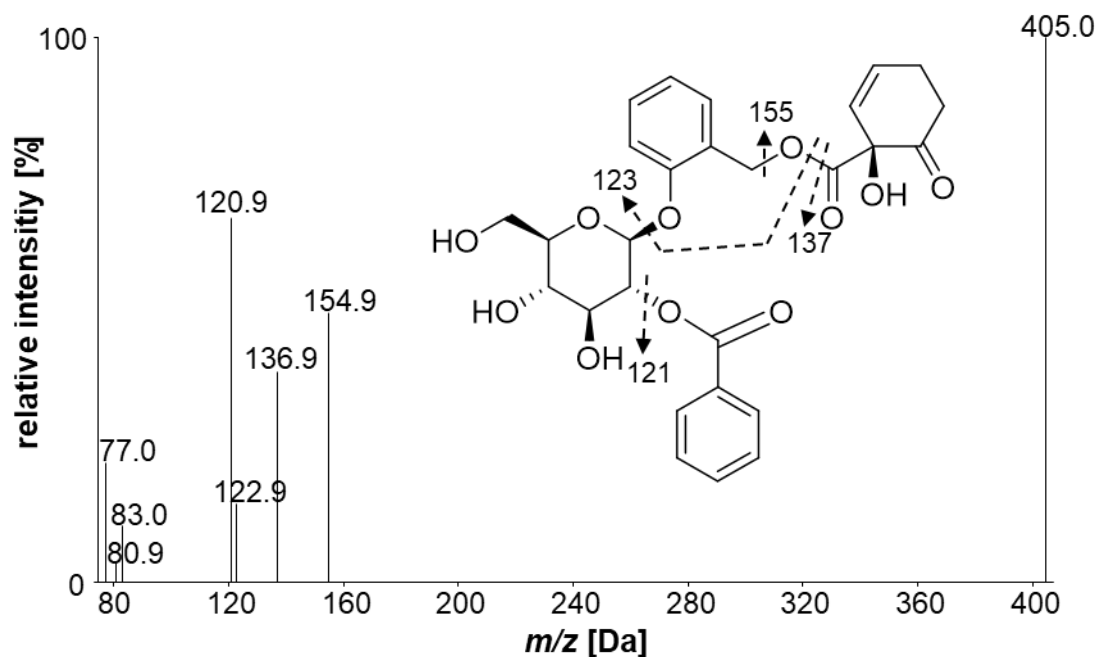


Figure S13: Centroided MS² spectrum of m/z 527.0 showing the fragmentation pattern of tremulacin (6).

5. Isolation and identification of compound 7 from subfraction F5-2

Fraction F5-2 (1 mg/mL in methanol/water (v/v 1/1)) was purified by means of semi-preparative UV-HPLC (λ = 252 nm, injection volume: 300 μ L, run time: 37 min) using the following chromatographic conditions: starting at 22% B, held there for 3 min, increased in 17 min to 45% B, held at 45% B for 2 min, increased further in 8 min to 50% B, held at 50% B for 3 min, decreased in 2 min to 22% B, and finally held isocratic for 2 min. Cinnamrutinose A was structurally elucidated in fraction 5-2-2 (7) using LC-ToF-MS and 1D/2D-NMR experiments.

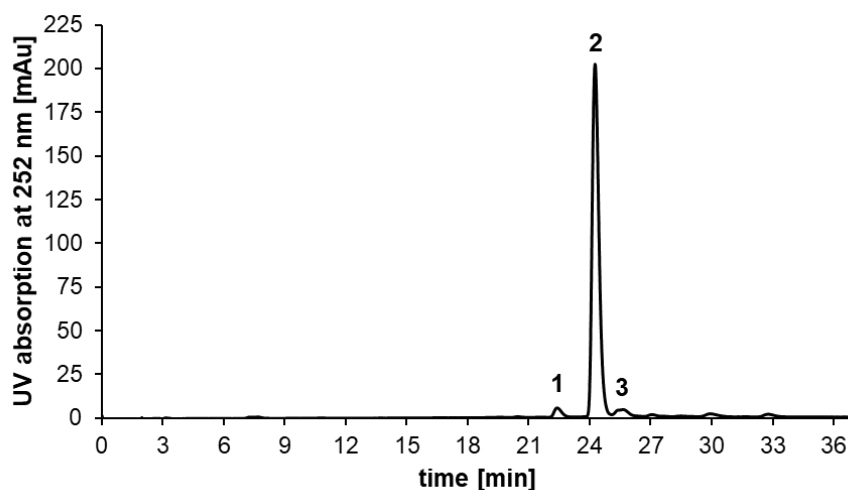


Figure S14: Semi-preparative UV-HPLC chromatogram of fraction F5-2 at 252 nm.

cinnamrutinose A (7)

$C_{21}H_{30}O_{10}$; UV (MeOH/H₂O, 1/1 v/v): λ_{\max} =204, 252 nm; LC-TOF-MS (ESI⁻): m/z 487.1824 [M+HCO₂H-H]⁻, 441.1768 [M-H]⁻; LC-MS/MS (DP = -55 V, CE = -56 V): m/z (%) 306.92 (100), 162.92 (41), 160.89 (3), 126.91 (1), 124.91 (14), 118.90 (13), 102.92 (19), 100.87 (7), 58.97 (4).

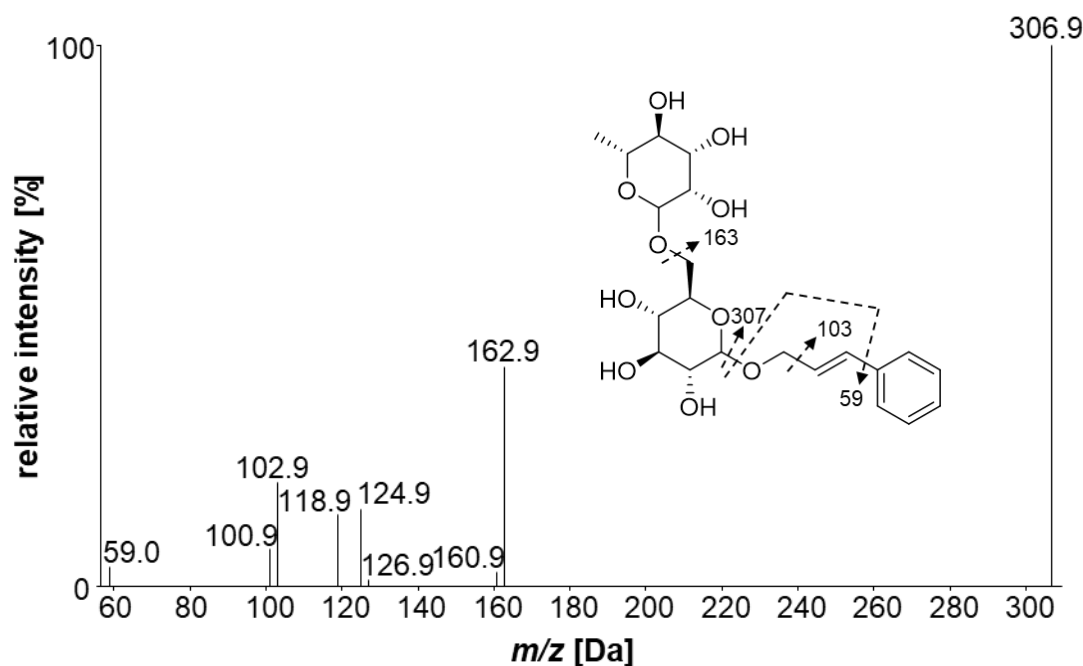


Figure S15: Centroided MS² spectrum of m/z 441.1 showing the fragmentation pattern of cinnamrutinose A (7).

6. QTRAP-LC-MS/MS conditions for sugar determination

For the determination of the sugar residues, the QTRAP-LC-MS/MS system was employed using 1% formic acid in H₂O (solvent A) and 1% formic acid in acetonitrile (solvent B) and positive ionization mode (ESI⁺). For chromatography a 2.1x100 mm, 1.7 μ m, Kinetex F5 column (Phenomenex, Aschaffenburg, Germany) was used and the following gradient (adapted from Schmid et al. (2018)) was applied: starting conditions were 5% of solvent B, held isocratic for 1.99 min at 5% B, then, increased to 20% in 3 min, in 21 min to 25% and in 1 min to 100 % B, held isocratic for 3 min at 100% B and decreased to 5% B in 1 min and finally held isocratic for 4 min at 5% B.

The MRM transitions Q1/Q3 of the derivatized sugars *D*-glucose (*m/z* 461.0/298.1), *L*-glucose (*m/z* 461.0/298.1), *D*-galactose (*m/z* 461.1/298.2), *D*-galacturonic acid (*m/z* 475.0/312.1), *D*-glucuronic acid (*m/z* 475.0/312.1), and *L*-rhamnose (*m/z* 445.0/282.1), as well as the DP, CE, and CXP values were adopted from Schmid et al. (2018).

7. Absolute configuration of salicylates

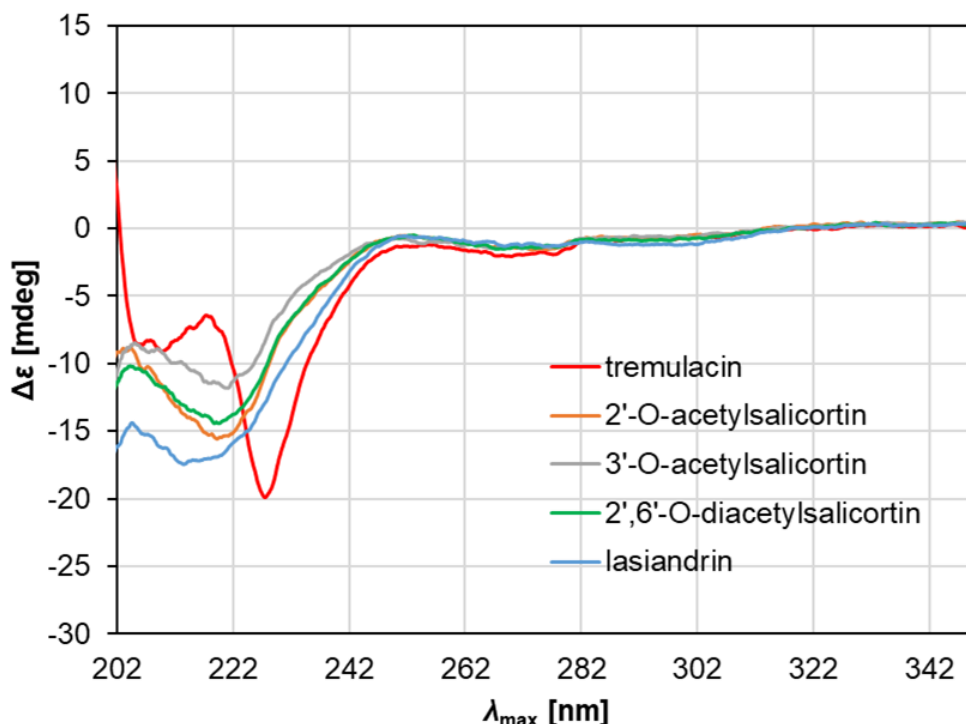


Figure S16: Absolute configuration of the isolated salicylates, 1-2 and 4-6, substituted with a HCH moiety determined by CD-spectroscopy. 2'-*O*-acetylsalicortin (1): $\Delta\epsilon = -15.5$ mdeg (*c* 0.43 mM, CH₃OH, $\lambda_{\max} = 220$ nm), 3'-*O*-acetylsalicortin (2): $\Delta\epsilon = -11.7$ mdeg (*c* 0.43 mM, CH₃OH, $\lambda_{\max} = 221$ nm), 2',6'-*O*-diacetylsalicortin (4): $\Delta\epsilon = -14.4$ mdeg (*c* 0.39 mM, CH₃OH, $\lambda_{\max} = 220$ nm), lasiandrin (5): $\Delta\epsilon = -17.2$ mdeg (*c* 0.33 mM, CH₃OH, $\lambda_{\max} = 216$ nm), tremulacin (6): $\Delta\epsilon = -19.6$ and -8.8 mdeg (*c* 0.38 mM, CH₃OH, $\lambda_{\max} = 228, 209$ nm).

8. Purification of fraction F7-4

For the semi-preparative UV-HPLC ($\lambda = 200$ nm) fractionation of F7-4 (15 mg/mL in methanol) a semi-preparative 250 × 10 mm, 5 μ m, PFP column was used. The chromatographic separation was performed using 0.1 % formic acid in water (solvent A) and 0.1 % formic acid in methanol (solvent B). The injection volume for both fractions was 100 μ L. The run time of the chromatographic separation of fraction F7-4 was 56 min. Chromatography of F7-4 started at 40% B, held at 40% B for 3 min, increased to 53% B in 17 min, held for 2 min, increased further to 57% B in 26 min, held for 2 min, and finally decreased again to 40% B in 3 min, which was held isocratic for 3 min. β -D-Glucopyranoside, 2-[[[(1-hydroxy-6,6dihydroxy-2-cyclohexen-1-yl)dihydroxy]oxy]methyl]phenyl, 2-acetate (8) was structurally elucidated in fraction F7-4-6 using LC-ToF-MS and 1D/2D-NMR experiments.

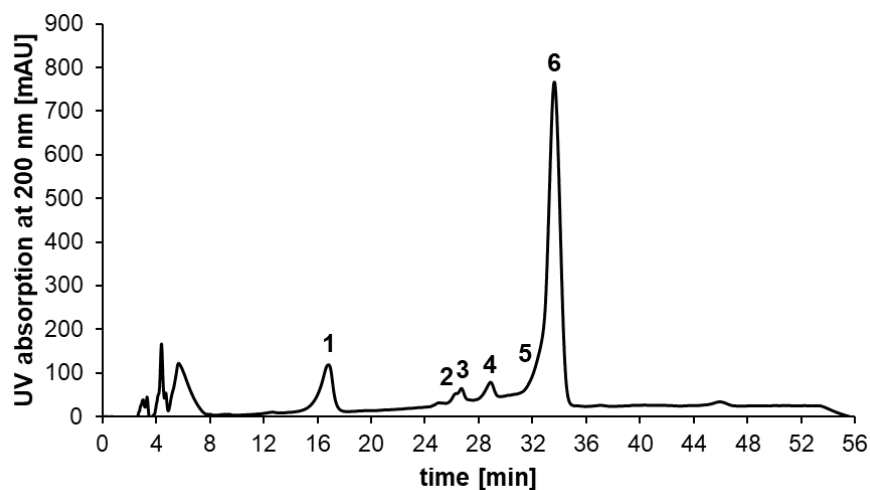


Figure S17: Semi-preparative UV-HPLC ($\lambda = 200$ nm) chromatogram of fraction F7-4.

β -D-Glucopyranoside, 2-[[[(1-hydroxy-6,6-dihydroxy-2-cyclohexen-1-yl)dihydroxy]oxy]methyl]phenyl, 2-acetate (8) in F7-4-6

$C_{22}H_{28}O_{12}$; LC-TOF-MS (ESI⁻): m/z 465.1401 [M-H₂O-H]⁻

After acetalization:

$C_{28}H_{36}O_{12}$; LC-TOF-MS (ESI⁻): m/z 563.2126 [M+two acetal groups-H]⁻

9. NMR spectroscopic data

The spectroscopic data of the isolated compounds are shown in Table 1-6. For NMR analysis, compound **6** was dissolved in DMSO-*d*₆, **1**, **2**, **4**, and **5** in acetone-*d*₆, **6** in acetonitrile-*d*₃, and **3** in methanol-*d*₄.

Table S1: ¹³C-NMR data of **1-6**. *compound 5: C(8) = [C=O], **compound 6: C(1'') = [C=O], C(2'') = [C].

Position	HSQC	δ_c [ppm]					
		1	2	3	4	5	6
1	[C]	155.88	156.41	155.85	155.78	155.85	154.20
2	[C]	125.94	125.83	131.92	126.10	126.17	124.34
3	[CH]	129.50	130.10	128.90	129.61	130.61	127.94
4	[CH]	123.36	123.10	123.70	123.56	123.66	122.30
5	[CH]	130.40	130.66	129.47	130.35	129.55	129.40
6	[CH]	116.49	116.35	115.96	116.62	116.80	115.05
7	[CH ₂]	63.08	63.67	59.97	63.07	62.17	61.40
8	[C=O]	170.79	170.86	-	170.78	170.78	169.86
9	[C]	78.81	78.83	-	78.85	78.86	77.35
10	[CH]	129.30	129.29	-	129.31	129.30	128.66
11	[CH]	132.43	132.44	-	132.41	132.43	131.56
12	[CH ₂]	27.20	27.17	-	27.20	27.20	25.85
13	[CH ₂]	36.09	36.11	-	36.06	36.07	35.56
14	[C=O]	206.20	206.21	-	206.20	206.18	205.92
1'	[CH]	100.18	102.03	100.75	100.06	100.25	98.41
2'	[CH]	74.39	72.87	75.07	74.26	74.41	73.79
3'	[CH]	75.66	77.70	75.98	75.06	75.45	74.29
4'	[CH]	71.44	69.36	71.41	71.47	71.39	69.89
5'	[CH]	78.05	78.46	78.35	75.50	75.05	77.28
6'	[CH ₂]	62.37	62.19	62.40	64.01	65.76	60.54
7'	[C=O]	-	-	-	170.92	-	-
8'	[CH ₃]	-	-	-	20.72	170.78*	-
9'	[C]	-	-	-	-	78.82	-
10'	[CH]	-	-	-	-	129.26	-
11'	[CH]	-	-	-	-	132.35	-
12'	[CH ₂]	-	-	-	-	27.20	-
13'	[CH ₂]	-	-	-	-	36.07	-
14'	[C=O]	-	-	-	-	206.18	-
1''	[CH ₃]	21.02	21.11	21.02	21.00	21.01	165.10**
2''	[C=O]	170.30	170.79	170.55	170.27	170.26	129.74**
3''	[CH]	-	-	-	-	-	129.34
4''	[CH]	-	-	-	-	-	128.68
5''	[CH]	-	-	-	-	-	133.38
6''	[CH]	-	-	-	-	-	128.68
7''	[CH]	-	-	-	-	-	129.34

Table S2: ¹H-NMR data of 1-6. *compound 5: C(8) = [C=O], ** compound 6: C(1'') = [C=O], C(2'') = [C].

Position	HSQC	δ_H [ppm]					
		1	2	3	4	5	6
1	[C]	-	-	-	-	-	-
2	[C]	-	-	-	-	-	-
3	[CH]	7.31	7.33	7.38	7.33	7.37	7.13
4	[CH]	7.05	7.05	7.03	7.06	7.08	7.00
5	[CH]	7.28	7.31	7.21	7.31	7.32	7.29
6	[CH]	7.21	7.25	7.13	7.22	7.22	7.18
7	[CH ₂]	5.18	5.23 5.35	4.55	5.13	5.18 5.12	4.80 4.95
8	[C=O]	-	-	-	-	-	-
9	[C]	-	-	-	-	-	-
10	[CH]	5.80	5.78	-	5.80	5.80	5.66
11	[CH]	6.14	6.13	-	6.14	6.14	6.07
12	[CH ₂]	2.49-2.54 2.67-2.76	2.48-2.53 2.64-2.71	-	2.48-2.53 2.66-2.75	2.46-2.54 2.62-2.73	2.39-2.46 2.47-2.56
13	[CH ₂]	2.59-2.66 2.94-2.86	2.54-2.57 2.83-2.90	-	2.54-2.58 2.85-2.93	2.53-2.59 2.84-2.92	2.47-2.52 2.69-2.63
14	[C=O]	-	-	-	-	-	-
1'	[CH]	5.13	5.11	5.05	5.16	5.17	5.29
2'	[CH]	5.02	3.64	5.03	5.06	5.02	5.05
3'	[CH]	3.72	5.08	3.66	3.82	3.74	3.69
4'	[CH]	3.57	3.66	3.50	3.58	3.52	3.34
5'	[CH]	3.58	3.63	3.51	3.77	3.85	3.51
6'	[CH ₂]	3.73 3.91	3.74 3.89	3.71 3.91	3.74 3.91	4.27 4.64	3.55 3.76
7'	[C=O]	-	-	-	-	-	-
8'	[CH ₃]	-	-	-	2.03	_*	-
9'	[C]	-	-	-	-	-	-
10'	[CH]	-	-	-	-	5.75	-
11'	[CH]	-	-	-	-	6.08	-
12'	[CH ₂]	-	-	-	-	2.46-2.54 2.62-2.73	-
13'	[CH ₂]	-	-	-	-	2.53-2.59 2.84-2.92	-
14'	[C=O]	-	-	-	-	-	-
1''	[CH ₃]	2.07	2.05	2.12	2.10	2.08	_**
2''	[C=O]	-	-	-	-	-	_**
3''	[CH]	-	-	-	-	-	7.98
4''	[CH]	-	-	-	-	-	7.52
5''	[CH]	-	-	-	-	-	7.65
6''	[CH]	-	-	-	-	-	7.52
7''	[CH]	-	-	-	-	-	7.98

Table S3: Multiplicity and coupling constants of **1-6**. *compound 5: C(8) = [C=O], ** compound 6: C(1'') = [C=O], C(2'') = [C].

Position	HSQC	Multiplicity; J [Hz]					
		1	2	3	4	5	6
1	[C]	-	-	-	-	-	-
2	[C]	-	-	-	-	-	-
3	[CH]	m	m	m	m	m	dd; 7.70, 1.51
4	[CH]	td; 7.50, 1.08	td; 7.50, 1.15	m	td; 7.44, 1.12	td; 7.42, 1.00	td; 7.62, 1.00
5	[CH]	m	m	m	m	dd; 7.78, 1.73	ddd; 7.97, 1.52
6	[CH]	d; 8.00	d; 7.84	dd; 8.20, 1.05	dd; 8.78, 1.06	d; 8.19	d; 8.21
7	[CH ₂]	d; 2.73 (overlap)	d; 12.80 d; 12.80	q; 15.46, 13.60 (overlap)	d; 13.07 (overlap)	d; 12.87 (overlap) d; 12.87	d; 13.37 d; 13.37
8	[C=O]	-	-	-	-	-	-
9	[C]	-	-	-	-	-	-
10	[CH]	dt; 9.70, 1.70	dt; 9.70, 1.70	-	dt; 9.81, 3.48	dt; 9.80, 1.70	dt; 9.90, 1.57
11	[CH]	dt; 9.70, 3.80	dt; 9.70, 3.80	-	dt; 9.67, 3.71	dt; 9.81, 3.96	dt; 9.81, 3.74
12	[CH ₂]	m m	m m	-	m m	m m	m m (overlap)
13	[CH ₂]	m m	m m	-	m m	m m	m (overlap) m (overlap)
14	[C=O]	-	-	-	-	-	-
1'	[CH]	d; 7.16	d; 7.70	d; 8.05	d; 7.16	d; 8.19	d; 8.13
2'	[CH]	dd; 8.08, 1.62	m	dd; 8.01, 1.46	dd; 8.33, 1.58	dd; 8.10, 1.55	dd; 8.08, 1.62
3'	[CH]	m	d; 9.30	m	m	t; 9.45	td; 5.49, 3.74
4'	[CH]	m	m	m	t; 9.31	t; 9.45	s
5'	[CH]	m	m	m	t; 9.44	ddd; 6.90, 5.12, 2.04	m
6'	[CH ₂]	m	dd; 11.35, 3.69 dd; 11.39, 2.99	m	dd; 11.75, 2.30 dd; 11.95, 6.24	dd; 11.80, 6.79 dd; 11.82, 2.06	m dd; 10.68, 4.99
7'	[C=O]	-	-	-	-	-	-
*8'	[CH ₂]	-	-	-	s	-	-
9'	[C]	-	-	-	-	-	-
10'	[CH]	-	-	-	-	dt; 9.80, 1.79	-
11'	[CH]	-	-	-	-	dt; 9.93, 3.89	-
12'	[CH ₂]	-	-	-	-	m m m	-
13'	[CH ₂]	-	-	-	-	m	-
14'	[C=O]	-	-	-	-	-	-
**1''	[CH ₂]	s	s	s	s	s	-
**2''	[C=O]	-	-	-	-	-	-
3''	[CH]	-	-	-	-	-	t; 7.94
4''	[CH]	-	-	-	-	-	m
5''	[CH]	-	-	-	-	-	m
6''	[CH]	-	-	-	-	-	t; 7.94
7''	[CH]	-	-	-	-	-	m

Table S4: NMR data (500/125 MHz) of **7**.

Position	HSQC	δ_c [ppm]	δ_H [ppm]	Multiplicity; J [Hz]
1	[CH ₂]	70.11	4.24 4.43	ddd; 12.83, 6.54, 1.44 ddd; 12.80, 7.04, 1.52
2	[CH]	126.84	6.35	ddd; 16.14, 5.61
3	[CH]	133.01	6.69	dt; 15.89, 1.82
4	[C]	137.80	-	-
5	[CH]	127.41	7.44	d; 7.60
6	[CH]	129.67	7.35	m
7	[CH]	128.73	7.30	m
8	[CH]	129.67	7.35	m
9	[CH]	127.41	7.44	d; 7.60
1'	[CH]	102.85	4.31	d; 7.75
2'	[CH]	74.70	3.14	t; 8.10, 8.59
3'	[CH]	71.25	3.22	m
4'	[CH]	77.63	3.37	m
5'	[CH]	73.75	3.25	m
6'	[CH ₂]	67.90	3.64 3.88	m dd; 11.46, 1.92
1''	[CH]	101.66	4.72	dd; 16.35, 1.03
2''	[CH]	71.74	3.78	dd; 3.78, 1.53
3''	[CH]	72.26	3.54	dd; 9.45, 3.63
4''	[CH]	76.34	3.29	m
5''	[CH]	69.07	3.59	m
6''	[CH ₃]	18.10	1.21	d; 6.41

Table S5: NMR data (500/125 MHz) of first diastereomer **8** (compound B) in methanol-*d*₄.

Position	HSQC	δ_c [ppm]	δ_H [ppm]	Multiplicity; <i>J</i> [Hz]
1	[C]	156.06	-	-
2	[C]	126.72	-	-
3	[CH]	129.67	7.39	m
4	[CH]	123.82	7.04	dt; 7.07, 1.14
5	[CH]	130.39	7.30	m
6	[CH]	116.85	7.21	ddd; 1.23, 7.57
7	[CH ₂]	63.13	5.11-5.18	m
8	[C=O]	171.58	-	-
9	[C]	86.75	-	-
10	[CH]	127.19	5.59	dd; 10.10, 2.75
11	[CH]	131.12	6.02	m
12	[CH ₂]	23.31	1.98-2.06	m
13	[CH ₂]	33.19	2.17-2.24	m
			1.93-2.00	m
			2.41-2.53	m
14	[C]	104.47	-	-
1'	[CH]	100.74	5.09	m
2'	[CH]	75.00	5.03	m
3'	[CH]	75.98	3.64-3.67	m
4'	[CH]	71.42	3.49	m
5'	[CH]	78.38	3.50	m
6'	[CH ₂]	62.41	3.75	m
1''	[CH ₃]	21.10	3.94	m
			2.15	s
2''	[C=O]	171.98	-	-

Table S6: NMR data (500/125 MHz) of second diastereomer **8** (compound C) in methanol-*d*₄.

Position	HSQC	δ_C [ppm]	δ_H [ppm]	Multiplicity; <i>J</i> [Hz]
1	[C]	156.06	-	-
2	[C]	126.72	-	-
3	[CH]	129.67	7.39	m
4	[CH]	123.82	7.04	dt; 7.07, 1.14
5	[CH]	130.39	7.30	m
6	[CH]	116.85	7.21	ddd; 1.23, 7.57
7	[CH ₂]	63.13	5.11-5.18	m
8	[C=O]	173.02	-	-
9	[C]	77.33	-	-
10	[CH]	128.44	5.53	dt (10.03, 2.01)
11	[CH]	132.80	5.98	dt (9.98, 3.62)
12	[CH ₂]	25.29	2.19-2.26	m
			2.33-3.39	m
13	[CH ₂]	32.13	2.44-2.51	m
14	[C]	111.81	-	-
1'	[CH]	100.74	5.09	m
2'	[CH]	75.00	5.03	m
3'	[CH]	75.98	3.64-3.67	m
4'	[CH]	71.42	3.49	m
5'	[CH]	78.38	3.50	m
6'	[CH ₂]	62.41	3.75	m
			3.94	m
1''	[CH ₃]	21.10	2.15	s
2''	[C=O]	171.98	-	-