

66. Sambucinic Acid, a New Metabolite of *Fusarium sambucinum*

45th Communication on Verrucarins and Roridins¹⁾

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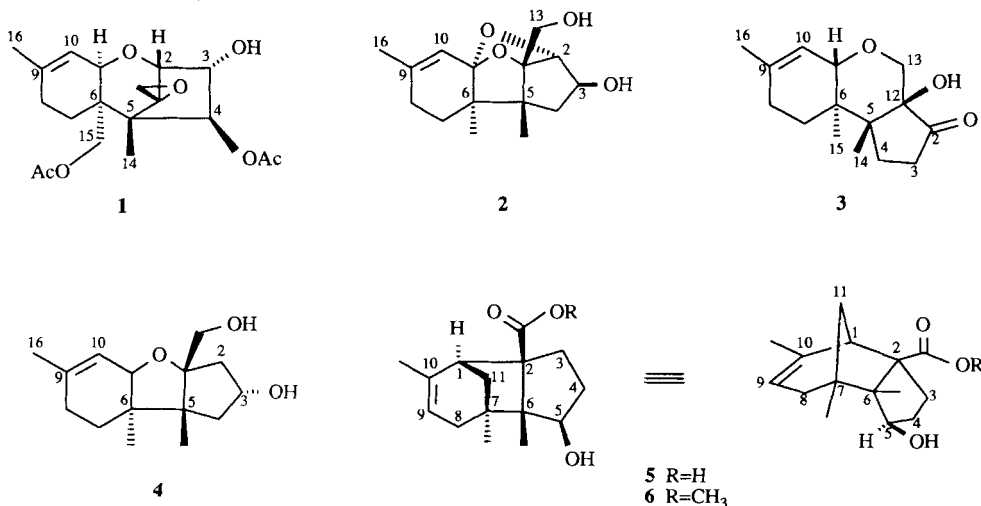
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Sambucinic acid (**5**), a C₁₅-trichothecene with a modified structure, has been isolated from cultures of *Fusarium sambucinum*. The structure is established by spectroscopic data and X-ray diffraction analysis. A hypothetical biogenetic pathway is proposed.

The trichothecenes are a still growing class of closely related sesquiterpenoid secondary metabolites produced by moulds, especially various species of *Fungi imperfecti* [2]. Many members of the family display a wide range of biological effects, such as cytostatic activity, but they are also highly toxic.

Recently, we reported on the structure elucidation of sambucinol (**2**) and sambucoin (**3**) which were isolated from cultures of *Fusarium sambucinum* (ATCC No. 11852) as minor metabolites [3]. The major compound was diacetoxyscirpenol (anguidine, **1**). In the



¹⁾ 44th Communication: [1].

Table 1. ^{13}C - and ^1H -NMR Data for Sambucinic Acid (**5**)^{a)}

Position	^{13}C Shift	^1H Shift	
1	46.76 (<i>d</i>)	2.69	(<i>d</i> , $J = 4$)
2	70.90 (<i>s</i>)	–	
3	28.19 (<i>t</i>)	1.72	(<i>td</i> , $J = 12, 6$)
		2.35	(<i>ddd</i> , $J = 12, 6.5, 2$)
4	38.47 (<i>t</i>)	2.27	(<i>ddd</i> , $J = 11, 5.5, 1.5$)
		2.67	(<i>m</i>)
5	75.70 (<i>d</i>)	4.44	(<i>dd</i> , $J = 10.5, 5.5$)
6	60.25 (<i>s</i>)	–	
7	42.54 (<i>s</i>)	–	
8	41.34 (<i>t</i>)	2.10	(<i>dquint.</i> , $J = 18, 2.5$)
		2.50	(<i>dm</i> , $J = 18$)
9	121.40 (<i>d</i>)	5.33	(<i>m</i>)
10	140.76 (<i>s</i>)	–	
11	46.11 (<i>t</i>)	1.82	(<i>d</i> , $J = 11$)
		2.90	(<i>ddd</i> , $J = 10.5, 4, 1.3$)
COOH	179.17 (<i>s</i>)	–	
C–C(6)	16.10 (<i>q</i>)	1.55	(<i>s</i>)
C–C(7)	25.41 (<i>q</i>)	1.15	(<i>s</i>)
C–C(10)	25.04 (<i>q</i>)	1.63	(<i>dt</i> , $J = 4, 2$)

^{a)} The spectra were recorded at 100.58 MHz (^{13}C) and 400 MHz (^1H) in (D_5)pyridine. Chemical shifts are given in ppm relative to TMS ($= 0.0$).

same large-scale fermentation, we have found a third minor metabolite, which we name sambucinic acid (**5**). Whereas diacetoxyscirpenol (**1**) possesses the normal trichothecane skeleton, the minor metabolites **2**, **3**, and **5** as well as 'apotrigothecene'²⁾ (**4**), a fourth minor metabolite isolated by a Canadian research group [5], are characterized by modified trichothecane skeletons.

Sambucinic Acid (= 5-Hydroxy-6,7,10-trimethyltricyclo[5.3.1.0^{2,6}]undec-9-ene-2-carboxylic acid; **5**). Yield ca. 0.4 mg/l. M.p. 200–207°. IR (KBr): 3430, 3400, 3200–2700, 1695. The composition of $\text{C}_{15}\text{H}_{22}\text{O}_3$ was secured by elemental analysis (found: C 70.36, H 9.29, calc.: C 72.00, H 8.80) and by EI-MS: 250 (M^+), 232, 205, 141. CI-MS (NH_3): 268 ($[M + \text{NH}_4]^+$), 250 (M^+), 233. MS-CI (CH_4): 251 ($[M + 1]^+$), 233, 189.

The structure elucidation was achieved by NMR spectroscopy, using modern pulse sequences³⁾. The ^1H - and ^{13}C -NMR assignments were made by a ^1H , ^{13}C heterocorrelation experiment (Table 1, Fig. 1).

There are signals of two olefinic C-atoms (C(9), C(10)), a secondary alcohol (C(5)), and a CO group at 179.17 ppm, which must be a carboxylic acid. Treatment of **5** with CH_2N_2 led to methyl sambucinate (**6**) (EI-MS: 264 (M^+), 205, 155 (100). CI-MS (NH_3): 282 ($[M + \text{NH}_4]^+$), 264 (M^+), 247 (100)). In the ^1H -NMR spectrum, a *singlet* at 3.67 ppm appeared, as expected.

A ^1H , ^1H -correlated 2D-NMR experiment (Fig. 2) with **5** clearly showed the following three isolated spin systems:

²⁾ The compound should correctly be named apotrigothec-9-ene-3 α ,13-diol, because apotrigothecane is the generic name for the skeleton [4].

³⁾ We thank Dr. Rügger, Spectrospin, Fällanden and Prof. H. Fritz, Ciba-Geigy AG, Basel, for recording the spectra and the helpful discussion.

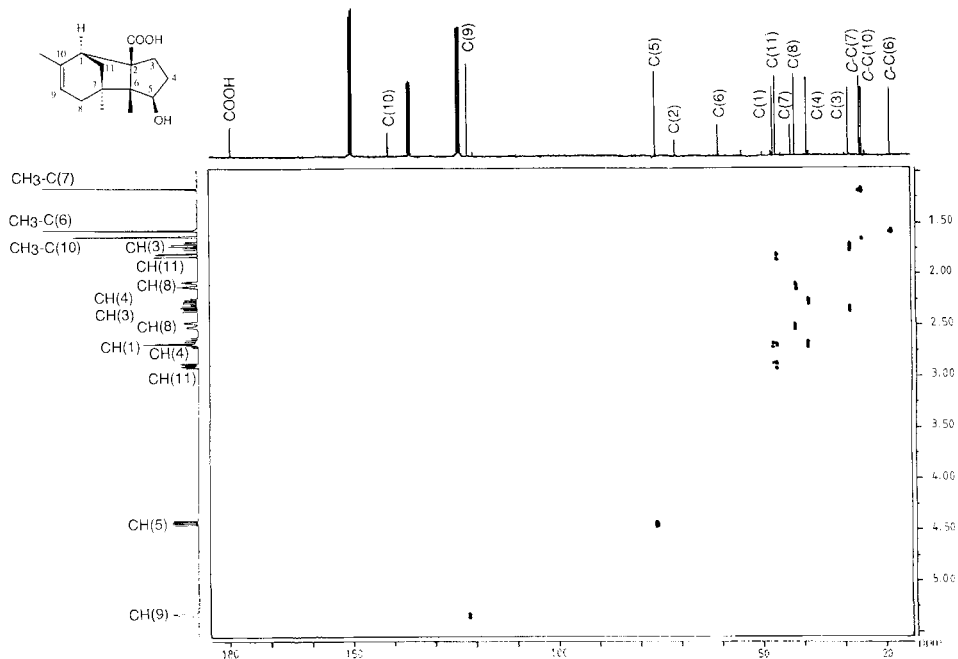


Fig. 1. ^{13}C , ^1H -Correlated 2D-NMR spectrum of sambucinic acid (5) (for conditions, cf. Table 1)

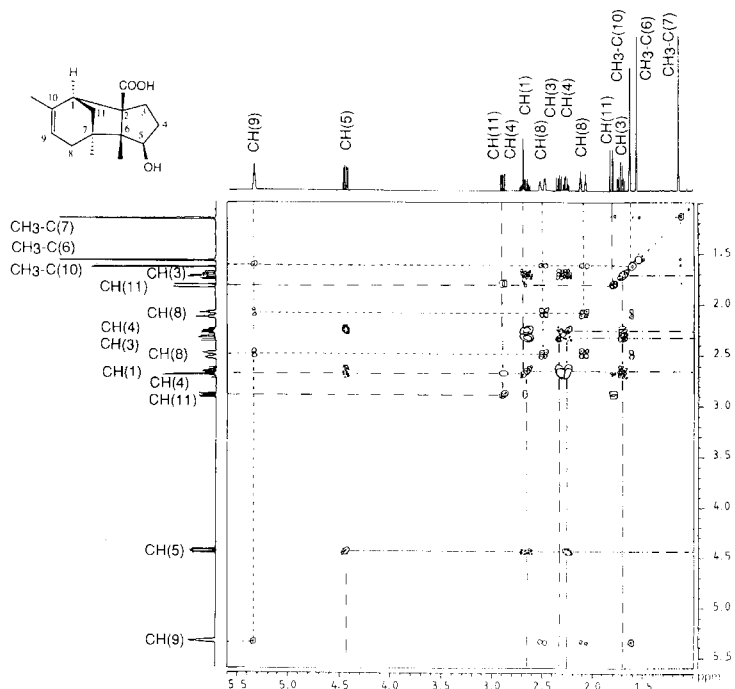


Fig. 2. ^1H , ^1H -Correlated 2D-NMR spectrum of sambucinic acid (5) (for conditions, cf. Table 1)

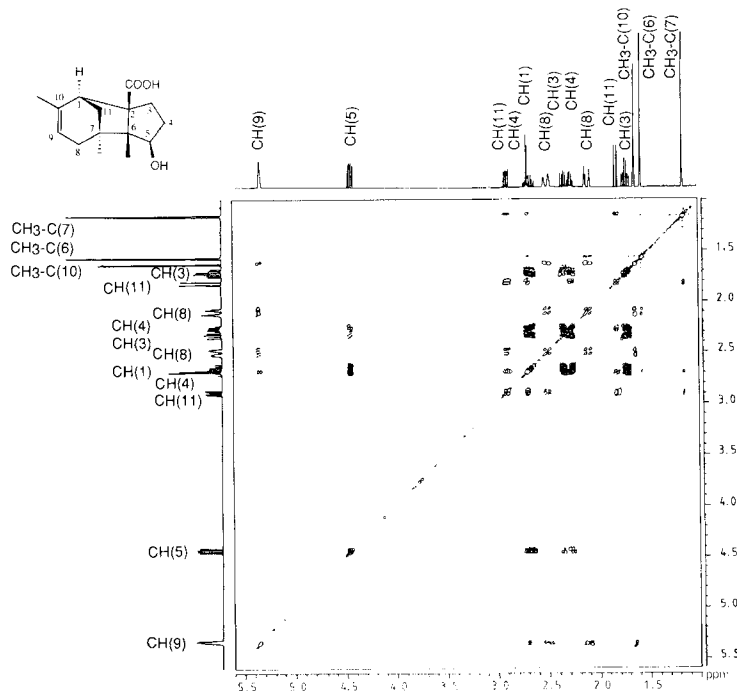


Fig. 3. $^1\text{H}, ^1\text{H}$ Long-range-correlated 2D-NMR spectrum of sambucinic acid (5) (for conditions, cf. Table 1)

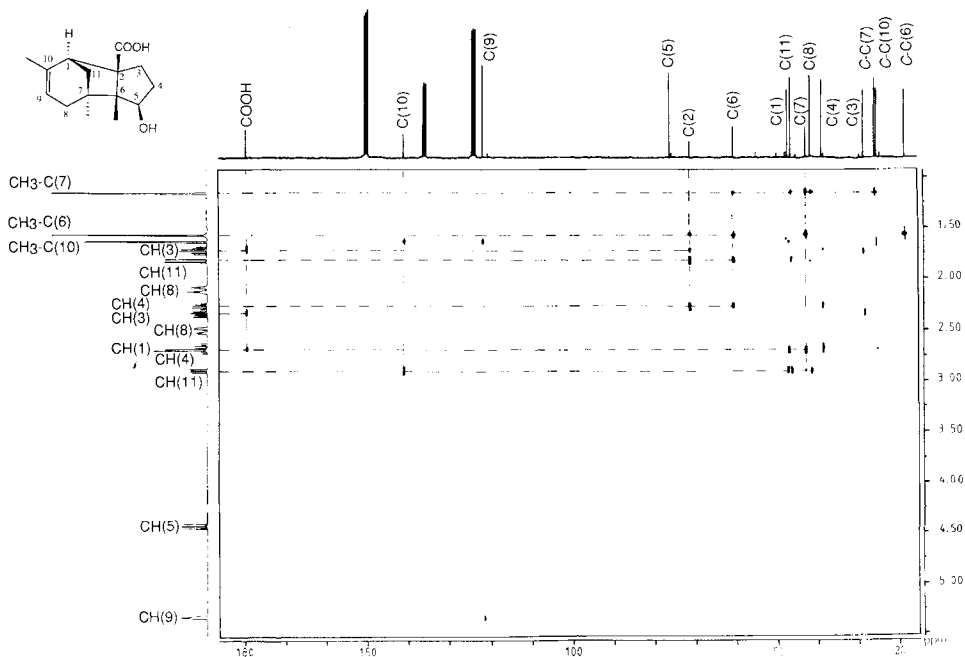
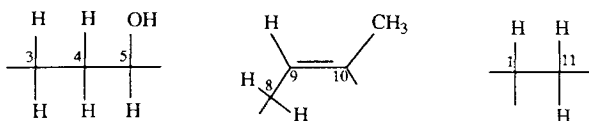


Fig. 4. $^{13}\text{C}, ^1\text{H}$ Long-range-correlated 2D-NMR spectrum of sambucinic acid (5) (for conditions, cf. Table 1)



The connection of the three spin systems and the location of the quaternary C-atoms and Me groups were established by a $^1\text{H}, ^1\text{H}$ long-range correlated 2D-NMR (Fig. 3) and a $^1\text{H}, ^{13}\text{C}$ heterocorrelation long-range 2D-NMR (Fig. 4): Long-range coupling of COOH with CH(1) and CH₂(3) clearly shows the sequence CH₂(3)–C(2)(COOH)–CH(1). C(2) (70.9 ppm) couples with CH(11), CH(3), CH(4), and CH₃–C(6). C(6) (60.2 ppm) shows long-range coupling with CH₃–C(6) (1.55 ppm), CH₃–C(7) (1.15 ppm), CH(4), and CH(11). C(7) (42.5 ppm) also couples with CH₃–C(6) and CH₃–C(7) and also with CH(1) and CH(11).

These facts and various additional correlations allowed to propose the structural formula for sambucinic acid.

The relative configuration and the location of CH₃–C(10) was assigned by a NOESY (Fig. 5) experiment. Hence, the configuration of C(5) was derived from a NOE of CH(5) with CH₃–C(7) and the attachment of CH₃–C(10) at the double-bond was determined by a NOE of CH₃–C(10) with CH(1).

The proposed structure was proven by X-ray diffraction analysis. Table 2 summarizes the crystal data, data collection details, and structure determination parameters. Unit-cell parameters were determined from accurate centering of 25 independent strong reflec-

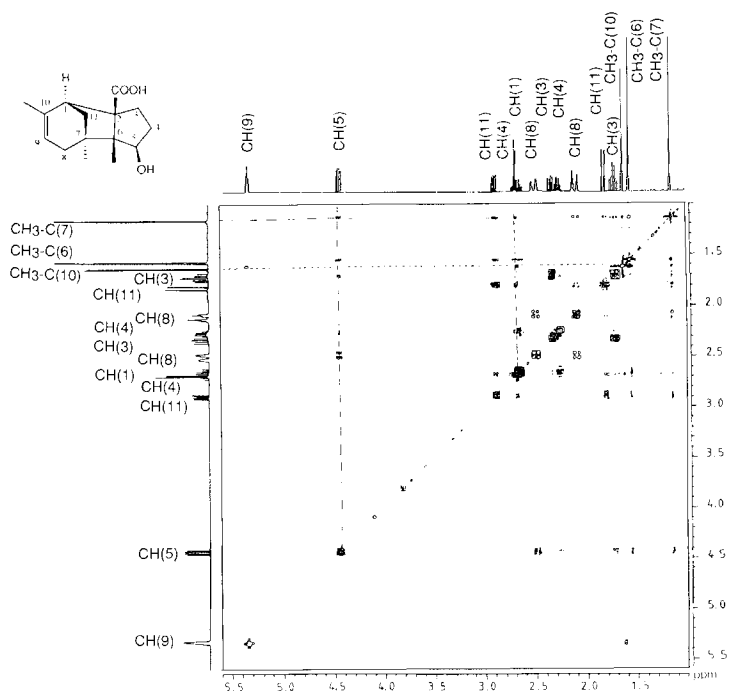


Fig. 5. NOESY Spectrum of sambucinic acid (5) (for conditions, cf. Table 1)

Table 2. *Crystal Data and Parameters of Data Collection for 5*

Formula	C ₁₅ H ₂₂ O ₃
Crystal system and space group	Orthorhombic, <i>P</i> 2 ₂ 1 ₂ 1 (No. 18 in non-standard setting)
<i>a</i> [Å]	7.793(2)
<i>b</i> [Å]	13.992(3)
<i>c</i> [Å]	24.409(6)
α [deg]	90.0
β [deg]	90.0
γ [deg]	90.0
<i>V</i> [Å ³]	2650.2
<i>Z</i>	8 (2 indep. molec./asymm. unit)
Crystal size [mm]	0.2 × 0.1 × 0.1
Temp. [K]	293
θ_{\max} [deg]	27
Radiation	MoK α ($\lambda = 0.71069$ Å)
Scan type	$\omega/2\theta$
No. of independent reflections	3322
No. of reflections used in refinements	1268 ($F > 2\sigma(F)$)
No. of variables	333
Final R_w	0.0770
Weighting system	$1.62/(\sigma^2(F) + 5.48 \cdot 10^{-4} F^2)$

tions by the least-squares method. Four standard reflections monitored every 3600 s during data collection showed no intensity loss. The raw data set was corrected for polarization effects. No correction for absorbance was applied. The final data set yielded a total of only 1268 independent reflections according to the criterion $F_o > 2\sigma(F_o)$. Crystals of C₁₅H₂₂O₃ have generally a weak intensity at every *Bragg* angle. Since there was no significant scattering observable in *h*00, one could not distinguish between the space groups *P*2₂1₂1 and *P*2₁2₁2₁. Density determination by flotation and the metric of the unit cell (18.4 Å³/non-H-atom) indicated two independent molecules per asymmetric unit. The

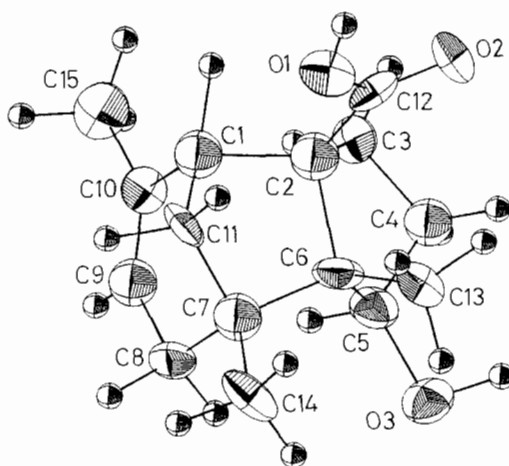
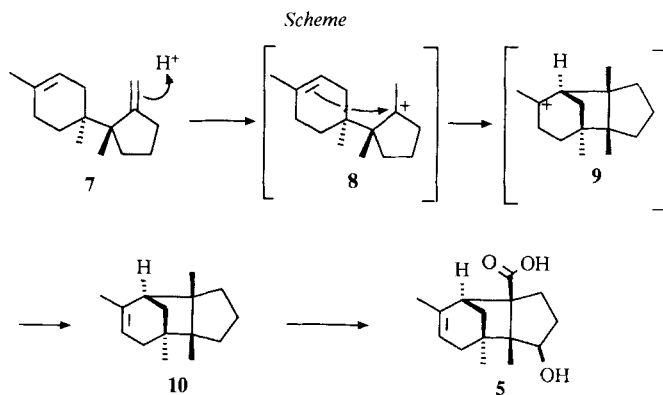


Fig. 6. ORTEP Plot of sambucinic acid (5) (arbitrary numbering)

programs MULTAN, MAGIC, YZARC led for both space groups to noninterpretable *E* maps, even when a large amount of starting phases were chosen. Finally, the structure was solved by the SHELXS-86 program [8] in *P*22₂₁ using direct methods strategies (TREF 180). Two independent molecules were localized per asymmetric unit.

Anisotropic least-squares refinements were made on 30 of the total 36 non-H-atoms; they were carried out on the SHELX-76 program [9]. H-atoms were partly localized from final ΔF maps and partly calculated. The refinements were stopped when Δ/σ was less than 0.1 for all non-H-atoms. Scattering factors for neutral atoms were taken from *Cromer et al.* [10], except those for H-atoms, which are from *Steward et al.* [11]. Fig. 6 shows an ORTEP plot of **5**. Fractional coordinates are deposited in the *Cambridge Crystallographic Data Base*. Structure factors, individual bond lengths, and bond angles are available from *M.Z.*

It is very likely that sambucinic acid (**5**) possesses the same absolute configuration as compounds **1-4**, because they are isolated from the same strain of the microorganism. Although the metabolites **1-5** exhibit different structures, they must be interrelated biogenetically. The biogenetic formation of sambucinol (**2**), sambucoin (**3**), and 'apotrichothecene' (**4**) can readily be rationalized assuming that they are derived from trichodiene (**7**) [3]. Compound **7** is a well-established intermediate in the biosynthesis of the normal trichothecenes [6][7]. It is reasonable to assume that also the biogenetic pathway of **5** branches off from **7** as shown in the *Scheme*:



Protonation of the exocyclic double bond of **7** leads to the carbenium ion **8**. By cyclization of the latter to cation **9** and subsequent stabilisation, the olefin **10** is formed. Hydroxylation at C(5) and oxidation will complete the formation of **5**.

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