

NARGENICIN BIOSYNTHESIS:
LATE STAGE OXIDATIONS AND
ABSOLUTE CONFIGURATION

DAVID E. CANE* and CHI-CHING YANG

Department of Chemistry, Brown University
Providence, Rhode Island 02912, U.S.A.

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Nargenicin A₁ (1)¹¹ and the closely related antibiotic nodusmicin (2)², are members of a novel class of saturated alicyclic polyketides containing a characteristic *cis*-fused octalin ring system. A similar, *trans*-fused, octalin nucleus is found in the Actinomycete metabolites chlorothricin (3)³, kijanimicin (4)⁴, and tetrocarcin A (5)^{5,6}, as well as in the fungal antibiotic ilicicolin H (6)⁷. Related to the latter metabolite are the recently described fungal phytotoxins betaenones A (7) and B (8)⁸ as well as the hypocholesterolemic agents compactin (9)⁹ and mevinolin (10)¹⁰. We have recently described incorporation experiments which establish the acetate and propionate origin of the carbon skeleton of nargenicin¹¹. Similar results were reported concurrently by RINEHART for the biosynthesis of nodusmicin¹². We also reported that incorporation of [1-¹⁸O₂, 1-¹³C]acetate gave nargenicin which bore oxygen-18 at C-1 and C-11, as detected by the isotopically induced shifts of the corresponding ¹³C NMR signals, while incorporation of [1-¹⁸O₂, 1-¹³C]propionate resulted in oxygen-18 enrichment at C-9 and C-17. These results have important implications for any proposed carbocyclization mechanism leading to the generation of the bicyclic octalin ring system from a reduced, linear polyketide precursor. The absence of propionate-derived oxygen at C-13 of nargenicin is inconsistent with a simple aldol-type condensation leading to the formation of the corresponding C-4, 13 bond, while the fact that neither the C-9 nor the C-11 oxygen atoms are derived from molecular oxygen rules out plausible epoxy-olefin or epoxy-alcohol cyclizations. As an alternative, we have suggested that the parent octalin ring system may be generated with the observed relative configuration and functional group distribution by an intramolecular Diels-Alder reaction between an *E,E*-4,6-diene and an *E* dienophile represented by the putative tetraene precursor

(11). Similar mechanisms may in fact be advanced to account for the formation of related octalin-containing metabolites*. In continuation of these studies, we have now determined the biosynthetic origins of the remaining oxygen atoms of nargenicin and established for the first time the absolute configuration of the nargenicin-nodusmicin family of antibiotics.

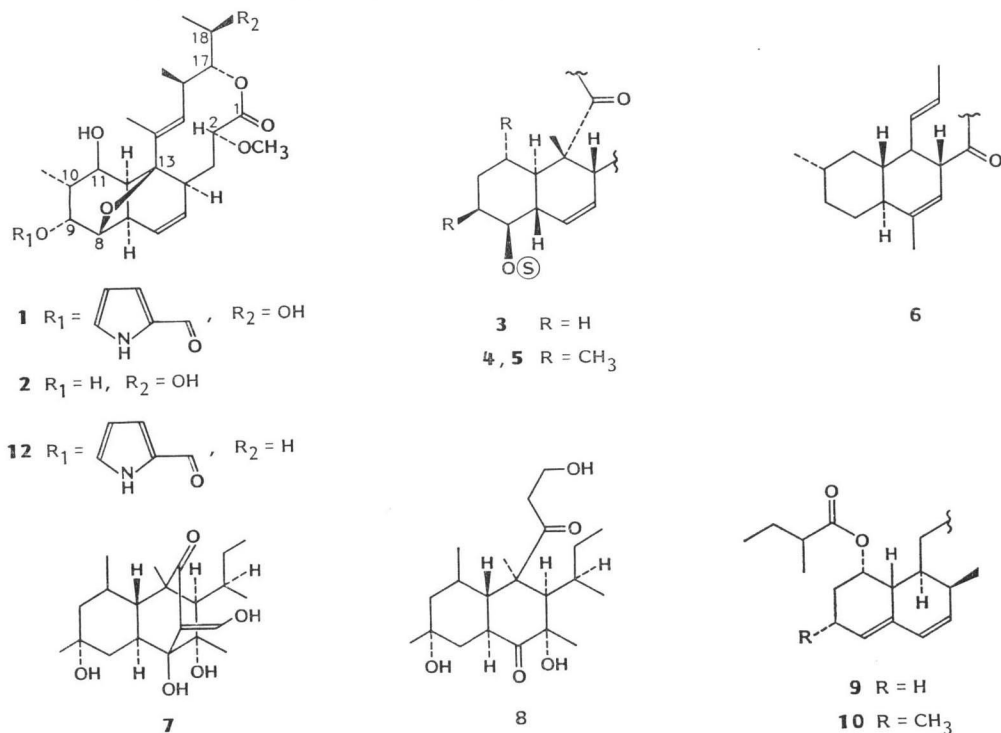
To determine the origin of the remaining oxygen atoms of nargenicin, *Nocardia argentinensis* Huang, ATCC 31306, was incubated in an atmosphere of ¹⁸O-labeled molecular oxygen. Considerable preliminary experimentation was necessary in order to establish suitable fermentation and labeling conditions due the severe reduction in nargenicin production associated with reduced culture aeration. A 2,800-ml Fernbach flask containing 280 ml of the previously described fermentation medium¹¹ was inoculated with 100-ml of a vegetative culture of *N. argentinensis* and incubated at 30°C and 250 rpm for 24 hours. The culture was then flushed with nitrogen and connected to a closed system containing a peristaltic pump and manostat which was used to circulate 4 liters of a 3:1 mixture of N₂ and ¹⁸O₂ (97 atom %) directly through the fermentation medium at a flow rate of 50~60 ml/minute**. A series of three U-tubes containing, respectively, charcoal, 5 N HCl and 5 N NaOH, was used to trap evolved CO₂ and other volatile materials. In order to increase the intensity of the relevant ¹³C NMR signals, and at the same time avoid unwanted excess intramolecular multiple ¹³C labeling, the cultures were supplemented with a mixture of 0.15 g of sodium [1-¹³C]propionate***, 0.15 g of sodium [2-¹³C]propionate***, 0.54 g of unlabeled sodium propionate, 0.18 g of sodium [2-¹³C]acetate***, 0.10 g of unlabeled sodium acetate, and 2.2 × 10⁷ dpm of sodium [1-¹⁴C]propionate at intervals of 24, 48 and 72 hours total incubation time in portions of 40%, 30% and 30%, respectively. After 96 hours, the resulting labeled nargenicin (10 mg) was isolated and purified as previously described¹¹. Analysis by high resolution 62.9 MHz ¹³C NMR revealed the pre-

* It is interesting to note that whereas the hypothetical *ring-forming* mechanism may be the same for nargenicin and for the *trans*-octalins 3, 4, and 5, the direction of polyketide *chain-elongation* is opposite for these two families of metabolites. See ref 15 and 16.

** For a description of this apparatus, see ref 17.

*** 90 Atom % ¹³C.

Fig. 1. Structures of saturated bicyclic polyketides, nargenicin **(1)**, nodusmicin **(2)**, betaenone A **(7)**, betaenone B **(8)** and 18-deoxynargenicin **(12)**, and partial structures of chlorothricin **(3)**, kijanimicin **(4)**, tetrocarcin **(5)**, ilicicolin H **(6)**, compactin **(9)** and mevinolin **(10)**.



Scheme 1. Proposed intramolecular Diels-Alder mechanism for biosynthesis of nargenicin and labeling of C-2, C-8, 13, and C-18 oxygen atoms by molecular oxygen.

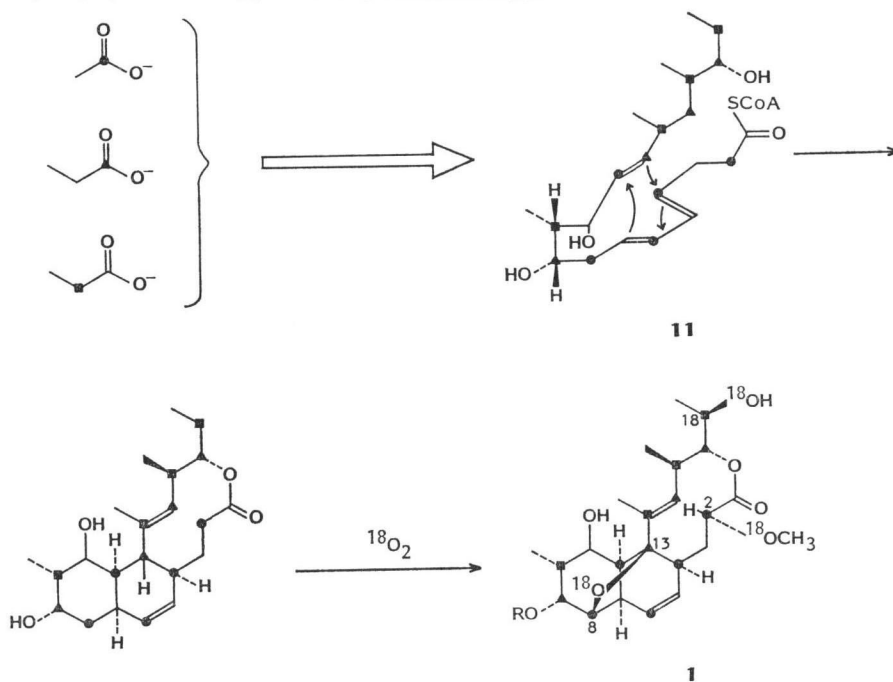


Table 1. Incorporation of $^{18}\text{O}_2$ into nargenicin A_1 ^a.

C	^{13}C shift (ppm)	Enrichment (%)	$\Delta\delta$ (ppm ^b)	$^{18}\text{O}:^{16}\text{O}^c$
2	83.4	4.9	0.018	35:65
8	81.3	4.9	0.024	30:70
13	89.1	3.0	0.030	30:70
18	66.6	8.9	0.024	25:75
23	57.6	1.6 ^d	0.024	20:80

^a Bruker WM 250, 62.9 MHz; spectral width 12195 Hz; 64 K data points; quadrature detection; 60° pulse; repetition time 1.34 seconds, LB=0 GB=0; 0.010 g sample in 0.4 ml of CDCl_3 .

^b $^{13}\text{C}^{18}\text{O}$ isotope shift, ± 0.006 ppm.

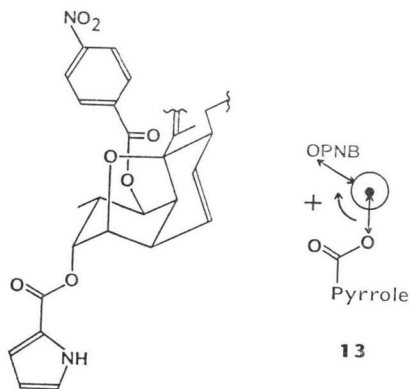
^c ± 5 ; uncorrected for contribution of natural abundance ^{13}C to $^{13}\text{C}^{18}\text{O}$ peaks.

^d Indirect ^{13}C labeling.

sence of isotopically shifted signals corresponding to excess ^{18}O at C-2, C-8, C-13, C-18, and the methoxyl methyl, C-23 (Table 1). These results complement the earlier data from incorporation of labeled acetate and propionate and establish unambiguously that the two ether oxygen atoms of nargenicin, as well as the C-18 hydroxyl oxygen are derived, as previously predicted¹¹, from molecular oxygen. Although it is not possible to stipulate the order in which the various oxygen atoms are introduced, it is likely that oxidation at C-18 occurs last, based on the reported isolation of 18-deoxynargenicin A_1 (**12**) from *Saccharopolyspora hirsuta*, the organism which produces nodusmicin¹³. To date, no further deoxygenated nargenicin or nodusmicin derivatives have been isolated and the detailed mechanism of formation of the C-8,13 ether bridge remains obscure.

While the structure and relative stereochemistry of nargenicin A_1 and nodusmicin have been unambiguously assigned by detailed ^1H NMR and X-ray crystallographic analyses^{1,2}, as well as by chemical conversion of **2** to **1**, to date the absolute configuration of these two antibiotics has not been reported. To solve this problem, we chose the nonempirical CD exciton chirality method of HARADA and NAKANISHI¹⁴. The ether-bridged, *cis*-fused octalin system provides a suitable rigid template bearing one of the two necessary chromophores, the pyrrole carboxylate moiety ($\lambda_{\text{max}}^{\text{MeOH}}$ 262 nm (ϵ 12,441)) which is esterified to the axial C-9 α hydroxyl group. Selective acetylation (1 equiv Ac_2O , pyridine,

Fig. 2. Partial structure of 11-*p*-nitrobenzoyl, 18-acetylnargenicin (**13**), showing clockwise relationship of pyrrole carboxylate and *p*-nitrobenzoate (PNB) chromophores.



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25°C, 18 hours) of the side chain (C-18) hydroxyl function, followed by *p*-nitrobenzoylation (*p*-nitrobenzoyl chloride, pyridine, catalytic 4-*N,N*-dimethylaminopyridine, reflux, 12 hours; λ_{max} 252 nm) of the equatorial C-11 β hydroxyl group, gave the corresponding 11-*p*-nitrobenzoyl, 18-acetylnargenicin (**13**), $\lambda_{\text{max}}^{\text{MeOH}}$ 260.5 (ϵ 15,036). The CD spectrum of **13** in MeOH showed a strong positive Cotton effect: $\Delta\epsilon$ (273.3 nm) +8.9, $\Delta\epsilon$ (252.3 nm) -2.4, $\Delta\Delta\epsilon$ 11.1. The positive Cotton effect corresponds to a clockwise relationship between the pyrrole carboxylate and *p*-nitrobenzoate chromophores¹⁴, as illustrated in Fig. 2. Nargenicin A_1 is therefore shown to have the absolute configuration represented by **1***.

Acknowledgment

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* The relative stereochemical assignments are based on the X-ray structure in Fig. 1 of ref 2. The usual way of drawing the configuration of the C-2 methoxyl group is ambiguous and does, in fact, tend to suggest the wrong relative stereochemistry at the latter center. Note also that the configuration at C-18 is incorrectly drawn in ref 11.

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