

Molecular Structure of the Metabolite Lanosulin

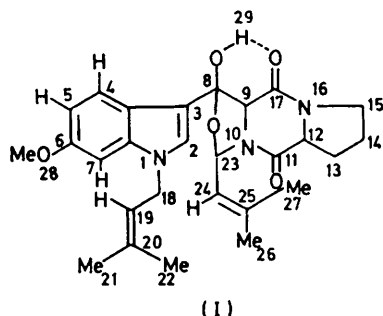
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Summary Structure (I) has been deduced for lanosulin the major metabolite of *Penicillium lanosum* Westling.

ECHINULIN (II),¹ neoehinulin (III),² brevianamides-A (IV),³ -B (V),⁴ and -E (VI),³ deoxybrevianamide-E (VII),^{3,5} and austamide (VIII)⁵ are a series of structurally similar metabolites which may be considered to be derived biogenetically from tryptophan, mevalonic acid, and alanine (II), glycine (III), or proline (IV)–(VIII). All formally possess an inverted $\gamma\gamma$ -dimethylallyl group at C-2, the origin of which has been discussed extensively, and several mechanisms have been proposed to account for its presence. Studies with model compounds tend to exclude both a direct S_N2' at the 2-position,⁶ or a primary attack at C-3 followed by rearrangement,⁷ but support a mechanism proceeding *via* primary attack at N-1 with subsequent rearrangement.⁸

We now report the structure of lanosulin (I) the major metabolite of *Penicillium lanosum* Westling. The $\gamma\gamma$ -dimethylallyl group at N-1 in lanosulin (I) presumably represents the missing biogenetic link for this group of compounds.

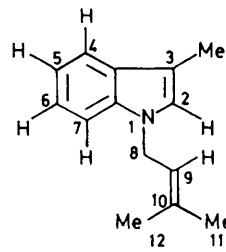


(I)

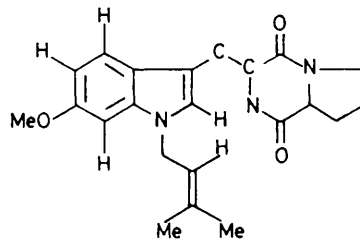
Lanosulin (I), $C_{27}H_{33}N_3O_5$, $[\alpha]_D^{25} + 24^\circ$ (c 0.91, $CHCl_3$), m.p. 204–206°, forms white crystals. Its u.v. spectrum† $[\lambda_{max}$ (EtOH) 228, 278, and 295 nm (ϵ 39,960, 11,000, and 12,290)] indicated the presence of an indole nucleus,⁹ and the n.m.r. data [δ 7.834 (4-H), 6.738 (5-H), and 6.650 (7-H) p.p.m.; $J_{4,7}$ 0.48, $J_{5,7}$ 2.15, $J_{4,5}$ 8.55 Hz] showed that a 5- or 6-methoxyindole was present. Comparison with the n.m.r. data for 2,3-dimethyl-5- and -6-methoxyindole suggested that the methoxy-group was at C-6, and this was confirmed by a Nuclear Overhauser effect (NOE) (6.6%) for 7-H on irradiation of the CH_2 part (18-H) of the $\gamma\gamma$ -dimethylallyl group at N-1. A similar enhancement for 5- and 7-H (Σ 19.9%) was observed on irradiation of 6-OMe. The 5- and 7-H resonances were also sharpened, indicative of a small coupling with 6-OMe.

The presence of the $\gamma\gamma$ -dimethylallyl group at N-1 was confirmed by comparison of the relevant n.m.r. data for (I) [δ ($CDCl_3$) 5.035 (19-H), 4.51 (18-H), 1.838 (22-Me), and 1.695 (21-Me) p.p.m.; $J_{18,19}$ 5.14, $J_{19,21-Me}$ 1.30 and $J_{19,22-Me}$ 0.96 Hz] with those for the indole (IX) [δ ($CDCl_3$) 5.34 (9-H), 4.53 (8-H), 1.74 (12-Me), and 1.69 (11-Me)

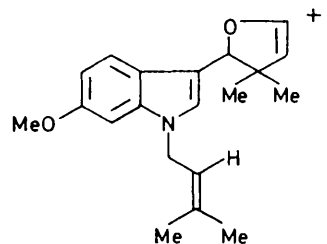
p.p.m.; $J_{8,9}$ 7.0, $J_{9,11-Me}$ 1.32, and $J_{9,12-Me}$ 1.0 Hz]. The NOE (19-H; 9.4%) detected on irradiation of 21-Me permits differentiation between the two methyl groups at C-21 and C-22.



(IX)



(X)



(XI)

The i.r. spectrum (Nujol) of (I) possesses two intense bands at 1685 and 1645 cm^{-1} which, together with the absence of amide-II bands suggested the presence of a piperazine-2,5-dione.³ Intramolecular hydrogen bonding¹⁰ [ν_{max} (CCl_4 ; 0.005M) 3500 cm^{-1} (OH)] is responsible for the low frequency of one of the carbonyl groups. Acid hydrolysis under standard peptide conditions gave proline. A prominent fragment in the mass spectrum at m/e 69 (C_6H_7N ; 62%) and n.m.r. data [δ 1.7–2.6 (13- and 14-H) and 3.533 (15-H) p.p.m.] provided further confirmation for the presence of a proline residue³ in (I).

Assuming that the indole fragment stems from tryptophan, these results indicate (X) as a part structure for lanosulin, leaving us to account for $C_6H_{10}O_2$. The strong intramolecular hydrogen bond in (I), and the n.m.r. and

† No change with acid or base.

high-resolution mass spectral results are consistent only with structure (I).

The base peak (XI) (m/e 311; $C_{20}H_{25}NO_2$)[†] stems from both m/e 479 ($C_{27}H_{33}N_3O_6$) and m/e 461 ($C_{27}H_{31}N_3O_4$) by loss of the piperazinedione fragment. Further, extensive fragmentation of the base peak proceeds along well defined pathways. The loss of HO and H_2O from (I) is readily accommodated. The C_5 unit attached to N-10 shows well defined n.m.r. resonances. Irradiation of 27-Me leads to a 11.6% enhancement of 23-H, and irradiation of 26-Me gives a 16.6% enhancement for 24-H.

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[†] Metastable transitions are available for all of these fragmentations.

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