

Methyl (1*S*,5*S*,6*S*,7*R*)-6-Formyl-7-methyl-2,8-dioxabicyclo[3.3.1]non-3-ene-4-carboxylate

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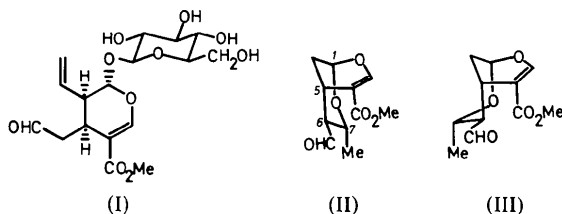
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Abstract. C₁₁H₁₄O₅, *M_r* = 226.23, orthorhombic, *P*2₁2₁2₁, *a* = 8.116 (4), *b* = 9.684 (4), *c* = 13.998 (6) Å, *U* = 1102 Å³, *Z* = 4, *D_x* = 1.364 Mg m⁻³, *μ*(Mo *Kα*) = 0.1 mm⁻¹. Final *R* = 0.040 for 1572 unique observed reflexions. The X-ray structure resolves the ambiguity of NMR measurements. The saturated and unsaturated rings adopt chair and sofa conformations respectively.

Introduction. The title compound was prepared from the natural product secologanin (I) *via* the action of β-glucosidase. Its proton NMR spectrum was consistent with either of two structures (II, III), which have different configurations at the C atoms bearing the methyl and formyl substituents; an X-ray structural investigation was undertaken to resolve this problem.



Prisms elongated along *a* were grown from diethyl ether/*n*-hexane. Data were collected to $2\theta_{\max} = 50^\circ$ on a Stoe STADI-2 diffractometer with monochromated Mo *Kα* radiation and two crystals mounted respectively about *a* (layers 0–7, 1466 reflexions) and *c* (layers 0–12, 1733 reflexions). After *L_p* corrections, averaging equivalent reflexions gave 1797 unique reflexions, 1573 with *F* > 4σ(*F*). Interlayer scale factors were obtained by least squares from equivalent reflexions from different layers; accurate cell constants were determined from ω angles of 279 strong reflexions.

The structure was solved by automated direct methods with *SHELXTL*; the best *E* map showed all

the non-hydrogen atoms. Refinement proceeded to *R* = 0.13 (isotropic), 0.10 (anisotropic). In the final cycles a riding model was employed for H atoms [C–H 0.96 Å, H–C–H 109.5°, *U*(H) = 1.2 *U*(C)]. The final *R'* = $\sum w^{1/2}\Delta / \sum w^{1/2}|F_o|$ was 0.046, with a corresponding *R* = 0.040. A final difference map showed no peaks

Table 1. Atom coordinates ($\times 10^4$) and isotropic temperature factors (Å² × 10³)

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> or <i>U_{eq}</i>
O(1)	5945 (2)	7141 (2)	6995 (1)	60 (1)*
C(2)	5384 (3)	8036 (3)	6328 (2)	47 (1)*
H(2)	4590	7699	5876	57
C(3)	5855 (3)	9359 (2)	6250 (2)	43 (1)*
C(4)	7071 (3)	9933 (3)	6950 (2)	47 (1)*
H(4)	6902	10912	6998	56
C(5)	6812 (4)	9171 (4)	7890 (2)	66 (1)*
H(5)	5709	9315	8116	77
H(5')	7581	9497	8360	77
C(6)	7096 (3)	7677 (3)	7697 (2)	61 (1)*
H(6)	6908	7210	8292	73
O(7)	8714 (2)	7400 (2)	7413 (1)	54 (1)*
C(8)	9251 (3)	8130 (2)	6565 (1)	40 (1)*
H(8)	8673	7829	6004	47
C(9)	8882 (3)	9670 (2)	6635 (2)	42 (1)*
H(9)	9604	10043	7112	50
C(10)	9208 (3)	10384 (2)	5704 (2)	53 (1)*
H(10)	9310	11372	5713	63
O(11)	9350 (3)	9816 (3)	4956 (1)	76 (1)*
C(12)	11065 (3)	7828 (3)	6476 (2)	59 (1)*
H(12)	11472	8283	5914	72
H(12')	11239	6851	6419	72
H(12'')	11642	8168	7027	72
C(13)	5180 (3)	10267 (2)	5512 (2)	45 (1)*
O(14)	5610 (3)	11447 (2)	5407 (2)	72 (1)*
O(15)	4018 (2)	9673 (2)	4973 (1)	59 (1)*
C(16)	3290 (4)	10529 (3)	4248 (2)	64 (1)*
H(16)	2519	9985	3888	76
H(16')	4114	10894	3826	76
H(16'')	2719	11276	4554	76

* *U_{eq}* is $\frac{1}{3}$ of the trace of the orthogonalized *U* matrix.

