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Key indicators

Single-crystal X-ray study T = 93 K Mean σ (C–C) = 0.003 Å R factor = 0.034 wR factor = 0.090 Data-to-parameter ratio = 12.9

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

15-(4-Chlorobenzylamino)-2-methoxypodocarpa-8,11,13-trien-15-one

At 93 K, the benzylic C–N bond of the title compound [systematic name: N-(4-chlorobenzyl)-6-methoxy-1,4a-dimethyl-1,2,3,4,4a,9,10,10a-octahydrophenanthrene-1-carboxamide], C₂₅H₃₀ClNO₂, adopts almost perfect coplanarity with the amide carbonyl linkage, the dihedral angle between the C–N–C=O planes being only 2.5 (3)°. There is weak intermolecular N–H···O hydrogen bonding between the two independent molecules comprising the asymmetric unit involving the carbonyl O atoms; this extends to link neighboring molecules into a chain.

Comment

As part of our anti-inflammatory and anticancer discovery program, we are exploring the derivatization of the C-4 carboxyl group of podocarpic acid (Bakare *et al.*, 2005) in order to design new molecules that can modulate the lipoxygenase and cyclooxygenase pathways. Consequently, the title compound, (I), was synthesized as one of a series of amide derivatives under investigation; its structure is reported here.



The geometric conformation parameters of the three fused six-membered rings in each of the two independent molecules comprising the crystallographic asymmetric unit of (I) (Fig. 1 and Table 1) are similar to those observed previously (Bakare et al., 2005; Couldwell et al., 1985; Mondal et al., 2003).In addition, the benzylic carbon-C19 to amide-N1 bond is almost coplanar with the amide carbonyl C15-O1 bond, with a dihedral angle of $2.5 (3)^\circ$. This near coplanarity is probably due to partial double-bond character of the C15-N1 bond with bond lengths of 1.348 (2) and 1.350 (2) Å for the two independent molecules. The major difference between the independent molecules is evident from Fig. 1, which shows a signifcant twist about the C4-C15 bond. There are weak intermolecular N-H···O hydrogen bonds between N and O1 in adjacent molecules, resulting in the formation of chains along the b axis (Fig. 2 and Table 2).

Experimental

To sodium hydride (NaH, 100 mg, 4.16 mmol) was added a solution of 12-methoxypodocarpa-8,11,13-trien-15-oic acid (537 mg, 1.86 mmol) in dry benzene (13 ml). The resulting mixture was stirred for 30 min,

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after which oxalyl chloride (2 ml) was slowly added and stirring continued for an additional 1 h. The reaction mixture was filtered and the filtrate concentrated *in vacuo* to give as a yellow oil, the acid chloride, 12-methoxypodocarpa-8,11,13-trienoyl chloride. Dry benzene (5 ml) was added to the residue and the solvent removed again *in vacuo*. The acid chloride, in benzene, was added slowly to a stirred solution of 4-chlorobenzylamine (in excess) in dry benzene (10 ml) at 273 K. The mixture was then allowed to warm to room temperature and stirred for 24 h before being filtered. The volatiles were removed *in vacuo* and the residue treated with hexane to give a white solid, which was purified by recrystallization from a solvent mixture of ethanol and distilled water (55%).

Crystal data

 $C_{25}H_{30}CINO_2$ $M_r = 411.95$ Triclinic, P1 a = 9.8466 (11) Å b = 10.2147 (10) Å c = 10.8492 (12) Å $\alpha = 84.224 (2)^{\circ}$ $\beta = 80.232 (2)^{\circ}$ $\gamma = 82.956 (2)^{\circ}$

Data collection

Bruker SMART 1K CCD diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996) $T_{\min} = 0.878, T_{\max} = 1$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.034$ $wR(F^2) = 0.090$ S = 1.046817 reflections 529 parameters H-atom parameters constrained $V = 1063.7 (2) \text{ Å}^{3}$ Z = 2 $D_{x} = 1.286 \text{ Mg m}^{-3}$ Mo K\alpha radiation $\mu = 0.20 \text{ mm}^{-1}$ T = 93 (2) KChunk, colorless $0.44 \times 0.33 \times 0.23 \text{ mm}$

8349 measured reflections 6817 independent reflections 6495 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.041$ $\theta_{\text{max}} = 28.2^{\circ}$

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0716P)^{2}]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} = 0.002$ $\Delta\rho_{max} = 0.37 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.37 \text{ e} \text{ Å}^{-3}$ Absolute structure: Flack (1983), 1588 Friedel pairs Flack parameter: 0.04 (4)

Table 1	I
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Selected geometric parameters (Å, °).

O2A-C12A	1.378 (3)	O2B-C12B	1.375 (2)
O2A-C18A	1.413 (3)	O2B-C18B	1.430 (2)
O1A-C15A	1.233 (2)	O1B-C15B	1.232 (2)
N1A-C15A	1.348 (2)	N1B-C15B	1.350 (2)
N1A-C19A	1.462 (3)	N1B-C19B	1.460 (2)
C4A-C15A	1.545 (3)	C4B-C15B	1.541 (2)
C4A-C16A	1.544 (3)	C4B-C16B	1.546 (3)
C4A - C5A	1.565 (2)	C4B-C5B	1.566 (2)
			()
C12A-O2A-C18A	117.61 (19)	C12B-O2B-C18B	116.89 (14)
C15A-N1A-C19A	120.34 (15)	C15B-N1B-C19B	122.46 (15)
C5A-C4A-C15A	117.98 (15)	C5B-C4B-C15B	113.92 (14)
C5A-C4A-C16A	109.55 (15)	C5B-C4B-C16B	109.45 (14)
O2A-C12A-C11A	115.13 (19)	O2B-C12B-C13B	124.18 (17)
O2A - C12A - C13A	124.42 (19)	O2B-C12B-C11B	115.91 (16)
O1A-C15A-N1A	120.03 (17)	O1B-C15B-N1B	120.97 (16)
O1A - C15A - C4A	121.72 (15)	O1B-C15B-C4B	120.05 (16)
N1A-C15A-C4A	118.14 (15)	N1B-C15B-C4B	118.83 (15)
N1A-C19A-C20A	113.14 (16)	N1B-C19B-C20B	111.61 (15)
			()
C19A-N1A-C15A-C	D1A 2.5 (3)	C19B-N1B-C15B-C	-1.2 (3)



Figure 1

View of the two independent molecules of (I) in the asymmetric unit, showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level.



Figure 2

The molecular packing of (I), viewed approximately down the *c* axis, showing the $N-H\cdots O$ hydrogen-bonding interactions (dashed lines) in the *b*-axis direction.

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1A - H1AC \cdots O1B$	0.88	2.19	2.9154 (19)	140
$N1B-H1BC\cdotsO1A^{i}$	0.88	2.23	2.9802 (19)	143
Symmetry code: (i) x y -	1 -			

Symmetry code: (i) x, y - 1, z.

The methyl H atoms were constrained to an ideal geometry, with C-H = 0.98 Å and $U_{iso}(H) = 1.5U_{eq}(C)$, but each group was allowed to rotate freely about its C-C bond. The position of the amine-H atoms were constrained to an ideal geometry, with N-H = 0.88 Å

and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm N})$. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H = 0.95-1.00 Å and $U_{\rm iso}({\rm H}) = 1.2U_{\rm eq}({\rm C})$.

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT-Plus* (Bruker, 1999); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1999); software used to prepare material for publication: *SHELXTL*.

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