Acta Crystallographica Section E Structure Reports Online

ISSN 1600-5368

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Key indicators

Single-crystal X-ray study T = 120 KMean σ (C–C) = 0.006 Å R factor = 0.091 wR factor = 0.256 Data-to-parameter ratio = 19.9

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

N-(3-Chlorophenyl)-2-(4-isobutylphenyl)propanamide

The asymmetric unit of the title compound, $C_{19}H_{22}ClNO$, contains two independent molecules with different conformations of the isobutyl group. The benzene rings in the two molecules make dihedral angles of 71.7 (1) and 76.6 (1)°. Intermolecular N-H···O hydrogen bonds link the molecules into infinite chains along [100].

Comment

Amides are a well known class of drugs having radio- and chemosensitizing properties (Pero et al., 1998) and are used in the treatment of psychiatric patients (Racagni et al., 2004). Iodobenzamide is the radiopharmaceutical imaging agent for the detection of melanoma and its metastases (Moins et al., 2002). Substituted benzamides are potential antipsychotic agents with antidopaminergic properties (Högberg et al., 1991) and inhibitors of steroid-5-reductase (Picard & Hartmann, 2002). Ibuprofen [2-(4-isobutylphenyl)propanoic acid], like naproxen, fenoprofen and benoxaprofen, is a classical nonsteroidal anti-inflammatory drug (NSAID) which inhibits the formation of prostaglandins, but a serious side effect is its gastrointestinal toxicity. A number of attempts to prepare derivatives with reduced side effects have been reported. Thus, ibuprofen sugar derivatives showed better anti-inflammatory activity than ibuprofen itself (Song et al., 2004). NSAIDs obtained by linking ibuprofen to furoxan and furazan groups exhibit anti-inflammatory activity comparable with that of ibuprofen, but with much reduced acute gastrotoxicity. The masking of the ibuprofen-free carboxylic group seems to be principally the basis of this reduced topical irritant action (Lolli et al., 2001). Ester and amide prodrugs of ibuprofen are significantly less irritating to the gastric mucosa than ibuprofen (Shanbhag et al., 1992). Amide derivatives of ibuprofen showed improved enantiomer separation using a chiral stationary phase in HPLC (Nicoll-Griffith, 1987). The title compound, (I), was prepared in the above context to obtain more effective NSAIDS with fewer side effects.



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The asymmetric unit of (I) contains two independent molecules (Fig. 1), having similar geometric parameters (Table 1), Received 23 August 2006 Accepted 30 August 2006





The asymmetric unit of the title compound, showing the atomic labelling and displacement ellipsoids drawn at the 50% probability level.

but differing in the orientation of the isobutyl group. In one molecule this group is positioned *trans* to the methyl group, while in the other molecule a *cis* arrangement is realized. The C112–C113–C116–C117 and C212–C213–C216–C217 torsion angles are -114.6 (5) and -107.9 (5)°, respectively. The mean planes of C101/N1/C107/O1/C108 and the chlorophenyl group make a dihedral angle of 32.6 (1)° [37.8 (1)° for the second molecule], and the two benzene rings make a dihedral angle of 71.7 (1)° [76.6 (1)° in the second molecule]. Intermolecular N–H···O hydrogen bonds (Table 2) link alternate independent molecules into infinite chains along [100], with the isobutylphenyl units oriented in parallel planes (Fig. 2).

Experimental

A solution of 2-(4-isobutylphenyl)propanoyl chloride (2.24 g, 10 mmol) in acetone (75 ml) was added dropwise to a solution of 3-bromoaniline (10 mmol) in acetone (10 ml) and the resulting mixture refluxed for 1 h. The reaction mixture was poured into cold water, whereupon the amide was precipitated as a solid. Repeated recrystallization attempts from ethanol provided colourless crystals (3.6 g, 8.1 mmol, 81%) of only very poor quality. The crystal finally selected for data collection was the best out of seven specimens investigated.

Crystal data

 $\begin{array}{l} C_{19}H_{22}CINO\\ M_r = 315.83\\ \text{Monoclinic, } P2_1/c\\ a = 9.8140 (16) \text{ Å}\\ b = 32.640 (5) \text{ Å}\\ c = 10.7522 (18) \text{ Å}\\ \beta = 91.276 (4)^\circ\\ V = 3443.3 (10) \text{ Å}^3 \end{array}$

Z = 8 D_x = 1.218 Mg m⁻³ Mo K α radiation μ = 0.22 mm⁻¹ T = 120 (2) K Prism, colourless 0.42 × 0.39 × 0.28 mm





The crystal packing, viewed down the c axis, with intermolecular hydrogen bonds indicated by dashed lines. H atoms not involved in hydrogen bonding have been omitted.

32978 measured reflections 7885 independent reflections

 $w = 1/[\sigma^2(F_0^2) + (0.1172P)^2]$

where $P = (F_0^2 + 2F_c^2)/3$

+ 1.5681P]

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 0.79 \ {\rm e} \ {\rm \AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.74 \text{ e} \text{ Å}^{-3}$

 $R_{\rm int} = 0.127$

 $\theta_{\rm max} = 27.5^{\circ}$

4890 reflections with $I > 2\sigma(I)$

Data collection

Bruker SMART CCD area-detector diffractometer φ and ω scans Absorption correction: multi-scan (*SADABS*; Bruker, 2002) $T_{min} = 0.922, T_{max} = 0.943$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.091$ $wR(F^2) = 0.256$ S = 1.097885 reflections 397 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

Cl1-C103	1.735 (4)	Cl2-C203	1.738 (4)
O1-C107	1.229 (4)	O2-C207	1.232 (4)
N1-C107	1.355 (5)	N2-C207	1.344 (5)
N1-C101	1.402 (5)	N2-C201	1.419 (5)
C107-C108	1.506 (5)	C207-C208	1.520 (5)
C108-C110	1.517 (5)	C208-C210	1.518 (5)
C108-C109	1.529 (5)	C208-C209	1.524 (6)
C107-N1-C101	126.4 (3)	C207-N2-C201	125.1 (3)
N1-C107-C108	115.7 (3)	N2-C207-C208	115.5 (3)
C107-C108-C110	109.6 (3)	C210-C208-C207	108.0 (3)

Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1 - H1A \cdots O2^{i}$ $N2 - H2A \cdots O1$	0.88 0.88	1.95 1.93	2.821 (4) 2.800 (4)	169 168
6	1.1			

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

H atoms were located in difference maps and refined in idealized positions riding on the C (C–H = 0.95–0.99 Å) or N (N–H = 0.88 Å) atoms, with $U_{\rm iso}({\rm H}) = 1.2 U_{\rm eq}({\rm C,N})$ and $1.5 U_{\rm eq}({\rm methyl}~{\rm C})$. Methyl H atoms were refined on the basis of rigid groups allowed to rotate but not tip.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Bruker, 2002); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

AS gratefully acknowledges a research grant from the Quaid-I-Azam University Islamabad Research Fund (project No. DFNS/ 2006-382).

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