Scattering factors from International Tables for Crystallography (Vol. C) Absolute structure: Flack (1983) Flack parameter = 0.0(2)

Table 1. Selected geometric parameters (Å, °)

S104	1.479 (5)	NI-C5	1.462 (7)
S1-C13	1.789 (6)	C2-C11	1.512 (8)
S1C12	1.802 (6)	C2—C3	1.544 (7)
O1C6	1.342 (6)	C3—C4	1.489 (8)
01-C7	1.463 (6)	C4—C5	1.517 (8)
O2—C6	1.212 (6)	C7-C10	1.457 (9)
03-C11	1.196 (7)	C7—C9	1.477 (9)
N1-C6	1.342 (7)	С7—С8	1.517 (8)
N1-C2	1.449 (7)	C11—C12	1.521 (8)
04	107.5 (3)	01C6N1	111.6 (4)
04-S1-C12	106.5 (3)	C10-C7-O1	110.7 (5)
C13—S1—C12	95.5 (3)	C10-C7-C9	112.9 (7)
C6-01-C7	121.3 (4)	O1—C7—C9	110.3 (5)
C6-N1-C2	124.8 (4)	C10-C7-C8	111.0 (7)
C6-N1-C5	121.6 (4)	O1-C7-C8	102.1 (4)
C2-N1-C5	113.6 (4)	C9—C7—C8	109.3 (5)
N1C2C11	112.5 (5)	O3-C11-C2	123.3 (5)
NI-C2-C3	102.2 (4)	O3-C11-C12	121.9 (6)
C11—C2—C3	111.1 (5)	C2-C11-C12	114.7 (5)
C4—C3—C2	103.7 (5)	C11-C12-S1	110.5 (4)
C3—C4—C5	105.5 (5)	C18C13C14	119.2 (6)
N1-C5-C4	102.7 (4)	C18-C13-S1	119.9 (5)
O2C6O1	125.7 (5)	C14-C13-S1	120.9 (5)
O2C6N1	122.7 (5)		

The proper enantiomer was chosen on the basis of the known configuration of the substrates and the method described by Flack (1983) was used to confirm the absolute configuration.

Data collection: XSCANS (Siemens, 1993). Cell refinement: XSCANS. Data reduction: XSCANS. Program(s) used to solve structure: SIR92 (Altomare et al., 1994). Program(s) used to refine structure: SHELXL97 (Sheldrick, 1997). Molecular graphics: SHELXTL/PC (Sheldrick, 1990). Software used to prepare material for publication: SHELXL97.

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Intermolecular N—H···N and C—H···O interactions form one-dimensional chains comprising the two independent molecules of N, N'-dicyclohexyl-N-nicotinovlurea

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Abstract

The title compound, C₁₉H₂₇N₃O₂, crystallizes in space group P1 with two molecules in the asymmetric unit which differ slightly in conformation. Intermolecular N-H···N and C-H···O interactions generate a hydrogen-bonded ring system between the alternating molecules, graph set $R_2^2(16)$, with N···N distances of 3.021 (3) and 3.041 (3) Å, and $C \cdots O$ distances of 3.219 (3) and 3.296 (3) Å along the hydrogen-bonded chains.

Comment

The general principles underlying molecular recognition processes are reasonably well understood and hydrogen bonding in crystal structures can usually be rationalized in preferred combinations of hydrogen-bond donors and acceptors (Etter et al., 1990). This allows comparison studies to be undertaken between classes of compounds containing analogous functional groups with a view to crystal engineering. However, compounds which are geometrically similar at the molecular level may differ at the supramolecular level, e.g. 2,2'-dipyridyl ketone and 2,2'-dipyridyl thioketone (Norsten et al., 1999). Thus, in molecules where several different potential hydrogenbond donors and acceptors are present (with cooperativity and/or competition among these interactions), the ability to deduce in advance the molecular packing arrangements in the crystal structure largely remains an unrealized vision (Wolff, 1996). The title compound, (I), a nicotinic acid derivative, forms part of a study of hydrogen-bonding interactions in a series of anion receptors (Gallagher & Fitzsimons, 1999; Gallagher *et al.*, 1999*a.b*).



Compound (I) crystallizes in space group $P\bar{1}$ with two independent molecules, A and B, in the asymmetric unit which differ slightly in conformation; views of the two molecules are depicted in Fig. 1. The r.m.s. deviation for the superposition of the non-H atoms in both molecules is 0.18 Å (Spek, 1998). Torsion-angle differences are evident for the C2--N1-C21-C22 angles, which are 87.0 (3) and 80.1 (3)° in molecules A and B, respectively (Table 2). Bond lengths and angles are unexceptional and in accord with anticipated values (Orpen et al., 1994), and selected dimensions are given in Table 1. The (sp^2) C1—C11 bond lengths of 1.498 (3) and 1.504 (3) Å are comparable with those in related structures (Orpen et al., 1994). The angles between the pyridinyl ring and the N1/C1/C2/C11 plane are 63.99 (12) and 58.62 (13)°; the angles between the four-carbon plane of the two cyclohexyl rings are 53.11 (12) and 65.71 (12)° in A and B, respectively.

Molecules A and B associate through intermolecular (amide)N—H···N(py) and (py)C—H···O—C(amide) interactions, generating hydrogen-bonded ring systems with graph set $R_2^2(16)$ between each pair (Table 2). Each molecule participates in hydrogen bonding both with N—H/C—H donors and (py)N/(amide)C—O acceptors, as depicted in the stereoview in Fig. 2. Chains are formed comprising alternating A and B molecules, along the a axis. Weak C—H···O—C intramolecular interactions are present involving cyclohexyl-ring-H atoms and the carbonyl-O atoms. The structure contains small voids in the crystal lattice of 7 Å³ (× 2), which are too small to accommodate a solvent molecule (*PLATON*; Spek, 1998).

Crystal structures with more than one molecule present in the asymmetric unit are not uncommon (Gallagher *et al.*, 1998) and we have reported previously structures with more than one molecule in the asymmetric unit in space groups $R\bar{3}$ (Ferguson *et al.*, 1992) and $P\bar{1}$ (Butler *et al.*, 1998). The rationalization of packing interactions in crystals with several independent mol-



Fig. 1. A view of (a) molecule A and (b) molecule B in (I) with the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

ecules is difficult (Karthe *et al.*, 1993). These may arise due to the existence of dimers or oligomers in solution (with energetically similar conformations arising from intramolecular hydrogen bonding, solute/solvent interactions *etc.*) and crystallizing to yield multiple formula units in the asymmetric unit (Desiraju, 1989). In (I), molecules A and B differ slightly in conformation and the overall crystal structure may be facilitated through hydrogen-bonded units crystallizing from solution to produce (I).

The crystal structure of N, N'-dicyclohexylurea has been reported (Govindasamy & Subramanian, 1997). A search of the Cambridge Structural Database (Allen & Kennard, 1993) for structures containing the 1,3-dicvclohexylurea moiety reveals several related compounds (Ball et al., 1990; Toniolo et al., 1990; Doucet et al., 1997).



Fig. 2. A stereoview of the intermolecular interactions along the $B \cdots A \cdots B \cdots A_{bottom}$ hydrogen-bonded chain; only the N-H and C---H H atoms participating in hydrogen bonding are included.

Experimental

The title compound was prepared by dissolving L-phenylalanine-L-leucine ethyl ester (1.63 g, 5 mmol) and nicotinic acid (0.62 g, 5 mmol) with 1-hydroxybenzotriazole (0.66 g, 5 mmol) in CH₂Cl₂ (50 ml). The mixture was cooled to 273 K and 1,3-dicyclohexylcarbodiimide (1.03 g, 5 mmol) was added. The reaction temperature was raised to room temperature after 30 min and stirred for 6 h. The title compound, (I), was isolated as the major product of the reaction by initial filtration of the reaction mixture and recrystallized from ethyl acetate/petroleum ether (313-333 K), yielding colourless crystals of (I).

Crystal data

$C_{19}H_{27}N_{3}O_{2}$	Mo $K\alpha$ radiation
$M_r = 329.44$	$\lambda = 0.7107 \text{ Å}$
Triclinic	Cell parameters from 25
PĪ	reflections
$a = 9.6855 (11) \text{\AA}$	$\theta = 5.44 - 19.44^{\circ}$
b = 12.775 (3) Å	$\mu = 0.079 \text{ mm}^{-1}$
c = 16.053 (6) Å	T = 290(1) K
$\alpha = 67.29 (3)^{\circ}$	Plate
$\beta = 87.640 (17)^{\circ}$	$0.44 \times 0.28 \times 0.14$ mm
$\gamma = 85.750 (17)^{\circ}$	Colourless
$V = 1827.0(8) \text{ Å}^3$	
Z = 4	
$D_x = 1.198 \text{ Mg m}^{-3}$	
D_m not measured	

Data collection

Enraf–Nonius CAD-4	$\theta_{\rm max} = 25.4^{\circ}$
diffractometer	$h = -11 \rightarrow 11$
ω -2 θ scans	$k = 0 \rightarrow 15$
Absorption correction: none	$l = -17 \rightarrow 19$
6786 measured reflections	3 standard reflections
6786 independent reflections	frequency: 240 min
2796 reflections with	intensity variation: <2%
$I > 2\sigma(I)$	2

Refinement

Refinement on F^2	$\Delta \rho_{\rm max} = 0.173 \ {\rm e} \ {\rm \AA}^{-3}$
$R[F^2 > 2\sigma(F^2)] = 0.047$	$\Delta \rho_{\rm min} = -0.143 \ {\rm e} \ {\rm \AA}^{-3}$
$wR(F^2) = 0.111$	Extinction correction:
S = 0.871	SHELXL97 (Sheldrick,
6786 reflections	1997a)
434 parameters	Extinction coefficient:
H atoms constrained	0.0181 (11)
$w = 1/[\sigma^2(F_o^2) + (0.0457P)^2]$	Scattering factors from
where $P = (F_o^2 + 2F_c^2)/3$	International Tables for
$(\Delta/\sigma)_{\rm max} = 0.001$	Crystallography (Vol. C

Table 1. Selected geometric parameters (Å, °)

		-	
01A—C1A	1.216(3)	O1 <i>B</i> —C1 <i>B</i>	1.216 (3)
02A—C2A	1.218 (3)	O2 <i>B</i> —C2 <i>B</i>	1.209 (3)
NIA-CIA	1.365 (3)	N1BC1B	1.362 (3)
N1A—C2A	1.445 (3)	N1 <i>B</i> —C2 <i>B</i>	1.444 (3)
N1A—C21A	1.480(3)	N1BC21B	1.483 (3)
N2A—C2A	1.323 (3)	N2B—C2B	1.333 (3)
N2AC31A	1.461 (3)	N2BC31B	1.453 (3)
C1A—C11A	1.498 (3)	C1 <i>B</i> —C11 <i>B</i>	1.504 (3)
C2A—N1A—C2	IA 117.5 (2)	C2B—N1B—C21B	116.8 (2)
C2A—N2A—C3	1A 123.2 (2)	C2BN2BC31B	121.6 (2)
01A-C1A-N1.	A 122.3 (2)	O1 <i>B</i> —C1 <i>B</i> —N1 <i>B</i>	122.8 (2)
01AC1AC1	IA 120.4 (2)	O1B—C1B—C11B	120.2 (2)
NIA-CIA-CI	IA 117.3 (2)	N1B—C1B—C11B	117.0 (2)
02A-C2A-N1	A 120.6 (2)	O2B—C2B—N1B	121.4 (2)
02A-C2A-N2	A 125.2 (2)	O2B—C2B—N2B	125.0(2)
N1A—C2A—N2	A 114.2 (2)	N2B—C2B—N1B	113.6 (2)
CIA	-NIA-C2A-02A	122.5 (3))
01 <i>A</i>		117.4 (3))
O1A	-C1A-C11A C16A	-58.8 (4)
C2A	-N1A-C21A-C22A	87.0 (3))
C2A	-N1A-C21A-C26A	-38.6 (3))
C1 <i>B</i> —N1 <i>B</i> —C2 <i>B</i> —O2 <i>B</i>		119.2 (3)	
O1B-C1B-C11B-C12B		123.3 (3)	
O1 <i>E</i>	C1B-C1B-C11B-C16B	-53.3 (4)
C2B	-N1B-C21B-C22B	80.1 (3))
C2 <i>B</i>	-N1B-C21B-C26B	-45.2 (3))

Table 2. Hydrogen-bonding geometry (Å, °)

D — $\mathbf{H} \cdots \mathbf{A}$	<i>D</i> —Н	H···A	$D \cdot \cdot \cdot A$	D — $H \cdot \cdot \cdot A$
N2A—H2A···N13B	0.86	2.21	3.041 (3)	163
C15A—H15A···O2B	0.93	2.50	3.296 (3)	143
$N2B - H2B \cdot \cdot \cdot N13A^{1}$	0.86	2.17	3.021 (3)	172
C15 <i>B</i> —H15 <i>B</i> ···O2A ¹	0.93	2.46	3.219 (3)	139
Symmetry code: (i) 1	+ x, y, z.			

Molecule (I) crystallized in the triclinic system, space group P1 or $P\overline{1}$; $P\overline{1}$ was assumed and confirmed by the analysis. H atoms were treated as riding atoms (C-H 0.93-0.98 and N—H 0.86 Å).

Data collection: CAD-4-PC Software (Enraf-Nonius, 1992). Cell refinement: SET4 and CELDIM in CAD-4-PC Software. Data reduction: DATRD2 in NRCVAX96 (Gabe et al., 1989). Program(s) used to solve structure: SHELXS97 (Sheldrick, 1997b). Program(s) used to refine structure: NRCVAX96 and SHELXL97 (Sheldrick, 1997a). Molecular graphics: NRC-VAX96, ORTEPII (Johnson, 1976) and PLATON (Spek, 1998). Software used to prepare material for publication: NRCVAX96, SHELXL97 and PREP8 (Ferguson, 1998).

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