## organic papers

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#### Key indicators

Single-crystal X-ray study T = 100 KMean  $\sigma$ (C–C) = 0.002 Å R factor = 0.045 wR factor = 0.116 Data-to-parameter ratio = 14.5

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

# N,N-Dimethyl-N'-(o-fluorobenzoyl)hydrazide

The title dimethylhydrazide o-FC<sub>6</sub>H<sub>4</sub>C(O)NHNMe<sub>2</sub>, or C<sub>9</sub>H<sub>11</sub>FN<sub>2</sub>O, is compared structurally both with other aroyl hydrazides, ArC(O)NHNH<sub>2</sub>, and with related trimethyl-ammonio ylides, Me<sub>3</sub>N(+)-N(-)C(O)Ar.

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## Comment

The title compound, (I), was prepared as a precursor to the corresponding trimethylammonio-stabilized nitrogen ylide  $Me_3N(+)-N(-)-C(O)C_6H_4$ -F-o. Its structure was determined in the course of an investigation into the ability of nitrogen ylides to act as ligands with transition metals (Morris *et al.*, 2003).



The presence of the *ortho*-fluoro substituent causes (I) to adopt a conformation (Fig. 1) which precludes conjugation across the C6–C7 bond [C1–C6–C7–N1 = 122.5 (2)°]. Possibly as a result, the C6–C7 bond [1.509 (2) Å] is 0.02 Å longer than the database average value obtained by Orpen *et al.* (1992) for such bonds, whereas all other bond lengths in (I) agree within 0.008 Å with appropriate average values from the Cambridge Structural Database (Allen, 2002). Phenylhydrazides, in general, adopt much flatter conformations than that found in (I). For example, in PhC(O)NHNH<sub>2</sub>, (II), the C(Ph)–C(Ph)–C(O)–N torsion angle is 30° (Kallel *et al.*, 1992), while in *o*-HOC<sub>6</sub>H<sub>4</sub>C(O)NHNH<sub>2</sub>, (III), the corresponding angle is only 3° (Mikenda *et al.*, 1993). In (I), the C6–C7–N1–N2 torsion angle is 175.9 (1)°. Corresponding values in other hydrazides are almost invariably close to 180°;



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View of (I), shown with 50% probability displacement ellipsoids.



#### Figure 2

Part of a stack of hydrogen-bonded molecules related by the *b*-glide plane normal to the a axis. The direction of view is down the a axis and b runs from left to right in the plane of the drawing.



The unit-cell contents, viewed in projection down the *a* axis.

the only exception we are aware of is provided by Ph(CO)NMeNMe<sub>2</sub>, where this torsion angle is only 8.5° (Knapp *et al.*, 1981), presumably for steric reasons. In (I), the N–N distance [1.420 (2) Å] and C–N–N angle [119.8 (1)°] are typical of hydrazides, such as (II) and (III), whereas in the closely related Me<sub>3</sub>N(+)–N(–)C(O)Ar ylides, the N–N distances are longer and the C–N–N angles more acute, *e.g.* 1.470 (3) Å and 114.2 (3)° when Ar = Ph (Cameron *et al.*, 1972), and 1.474 (1) Å and 114.8 (1)° when Ar = *p*-Cl–C<sub>6</sub>H<sub>4</sub> (Morris *et al.*, 2003). *QUEST* and *CONQUEST* search programs were used with Version 5.24 of the Cambridge Structural Database (Allen, 2002) to obtain some torsion angles not given by the original authors.

Glide-plane-related molecules connected by  $N-H\cdots O$ hydrogen bonds (Fig. 2) form stacks which run parallel to the *b* axis. These stacks pack so that alternate layers parallel to (001) contain, respectively, aromatic rings and terminal  $NMe_2$  groups, as can be seen by viewing the contents of the unit cell in projection down the *a* axis (Fig. 3). As expected (Glusker *et al.*, 1994), the F atoms do not participate in the hydrogen bonding.

The atomic  $U^{ij}$  values are moderately well reproduced by a **TLS** analysis (Schomaker & Trueblood, 1968):  $R_2 = (\Delta U^2/U^2)^{1/2} = 0.10$ ; they also pass the Hirshfeld (1976) rigid-bond test, the worst discrepancy being  $\Delta U = 0.0018 (12) \text{ Å}^2$  for C4–C5.

## Experimental

The title compound was prepared by the method of Smith *et al.* (1968).

 $R_{\rm int} = 0.057$ 

 $\theta_{\rm max} = 25.9^{\circ}$ 

 $h = -14 \rightarrow 14$  $k = -9 \rightarrow 9$ 

 $l = -24 \rightarrow 24$ 

+ 0.85P]

 $\Delta \rho_{\rm min} = -0.25 \text{ e} \text{ Å}^{-3}$ 

 $(\Delta/\sigma)_{\rm max} = 0$  $\Delta\rho_{\rm max} = 0.23 \text{ e} \text{ Å}^{-3}$ 

 $w = 1/[\sigma^2(F_o^2) + (0.048P)^2]$ 

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

Crystal data

C<sub>9</sub>H<sub>11</sub>FN<sub>2</sub>O Mo  $K\alpha$  radiation  $M_r = 182.2$ Cell parameters from 1794 Orthorhombic, Pbca reflections a = 11.8503 (5) Å $\theta = 2.9 - 26.0^{\circ}$  $\mu = 0.10 \text{ mm}^{-1}$ b = 7.8530(3) Å c = 19.9630(9) Å T = 100 (2) K $V = 1857.77 (13) \text{ Å}^3$ Needle, colourless Z = 8 $0.40 \times 0.05 \times 0.05 \text{ mm}$  $D_x = 1.303 \text{ Mg m}^{-3}$ 

#### Data collection

Nonius KappaCCD diffractometer Thick-slice  $\omega$  and  $\varphi$  scans 6796 measured reflections 1797 independent reflections 1491 reflections with  $I > 2\sigma(I)$ 

Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.045$   $wR(F^2) = 0.116$  S = 1.111797 reflections 124 parameters H atoms treated by a mixture of independent and constrained refinement

#### Table 1

Hydrogen-bonding geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1-H1\cdotsO1^{i}$	0.91 (2)	1.98 (2)	2.8801 (19)	168.8 (19)
Summatry and (i) 1 x y 1 z				

Symmetry code: (i)  $\frac{1}{2} - x$ ,  $y - \frac{1}{2}$ , z.

All H atoms were initially located from difference syntheses. The H atom bonded to N1 was freely refined. Other H atoms were positioned geometrically and refined with riding constraints, assuming C–H bond lengths of 0.93 and 0.96 Å, respectively, for  $sp^2$  and  $sp^3$  C atoms. In the case of each of the two methyl groups, a single orientation parameter was also refined.

Data collection: COLLECT in KappaCCD Control Software (Nonius, 2000); cell refinement: HKL SCALEPACK (Otwinowski & Minor, 1997); data reduction: HKL DENZO (Otwinowski & Minor, 1997) and SCALEPACK; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEP-3 for Windows (Farrugia, 1997); software used to prepare material for publication: WinGX (Farrugia, 1999).

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