

N-Isopropyl-3-methyl-2-nitrobenzamide

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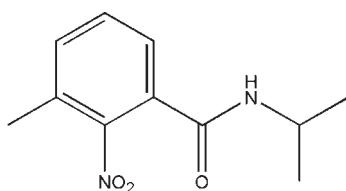
Received 3 July 2009; accepted 12 August 2009

Key indicators: single-crystal X-ray study; $T = 298\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.006\text{ \AA}$; R factor = 0.066; wR factor = 0.174; data-to-parameter ratio = 15.5.

In the title compound, $\text{C}_{11}\text{H}_{14}\text{N}_2\text{O}_3$, the bond lengths and angles are within normal ranges. Weak intermolecular $\text{N}-\text{H}\cdots\text{O}$ interactions link the molecules into chains along the a axis. A non-classical intramolecular $\text{C}-\text{H}\cdots\text{O}$ interaction (nitro O atom and a H atom of the nearest methyl group) is found, forming a six-membered ring with a twisted conformation. This six-membered ring has a twisted conformation.

Related literature

For bond-length data, see: Allen *et al.* (1987). For general background, see: Lahm *et al.* (2005).

**Experimental***Crystal data*

$\text{C}_{11}\text{H}_{14}\text{N}_2\text{O}_3$

$M_r = 222.24$

Orthorhombic, $Pbca$

$a = 9.4230 (19)\text{ \AA}$

$b = 13.250 (3)\text{ \AA}$

$c = 20.041 (4)\text{ \AA}$

$V = 2502.2 (9)\text{ \AA}^3$

$Z = 8$

Mo $K\alpha$ radiation
 $\mu = 0.09\text{ mm}^{-1}$

$T = 298\text{ K}$
 $0.30 \times 0.20 \times 0.10\text{ mm}$

Data collection

Enraf–Nonius CAD-4
diffractometer
Absorption correction: ψ scan
(North *et al.*, 1968)
 $T_{\min} = 0.974$, $T_{\max} = 0.991$
2260 measured reflections

2260 independent reflections
1135 reflections with $I > 2\sigma(I)$
3 standard reflections
every 200 reflections
intensity decay: 1%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.066$
 $wR(F^2) = 0.174$
 $S = 1.00$
2260 reflections

146 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.22\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.16\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N}1-\text{H}1\text{A}\cdots\text{O}1^{\text{i}}$	0.86	2.00	2.855 (3)	173
$\text{C}11-\text{H}11\text{A}\cdots\text{O}3$	0.96	2.37	3.021 (5)	124

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

Data collection: *CAD-4 EXPRESS* (Enraf–Nonius, 1994); cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

The authors thank the Center of Testing and Analysis, Nanjing University, for support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RK2153).

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supporting information

Acta Cryst. (2009). E65, o2278 [doi:10.1107/S1600536809031997]

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S1. Comment

The title compound contains nitro- and acetyl amino-groups, which can react with different groups to prepare various functional organic compounds as a fine organic intermediate (Lahm *et al.*, 2005). We herein report the crystal structure.

In the title molecule (**I**), (Fig. 1), the bond lengths and angles are within normal ranges (Allen *et al.*, 1987).

Intramolecular C—H···O interaction (Table 1) results in the formation of a six-membered ring (O3/N2/C10/C9/C11/H11A), having twisted conformation.

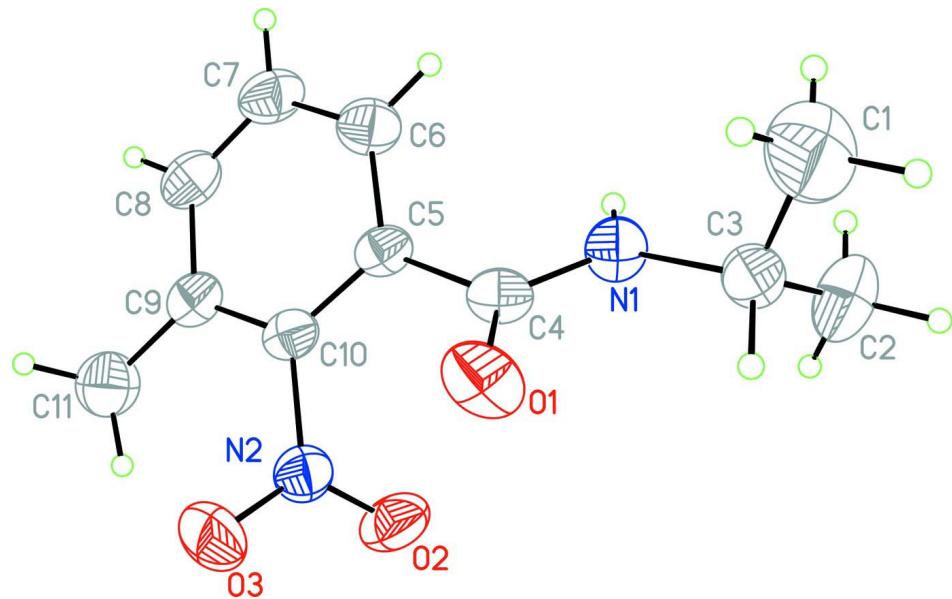
In the crystal structure, weak intermolecular N—H···O interactions (Table 1) link the molecules into chains along the *a* axis (Fig. 2), in which they may be effective in the stabilization of the structure.

S2. Experimental

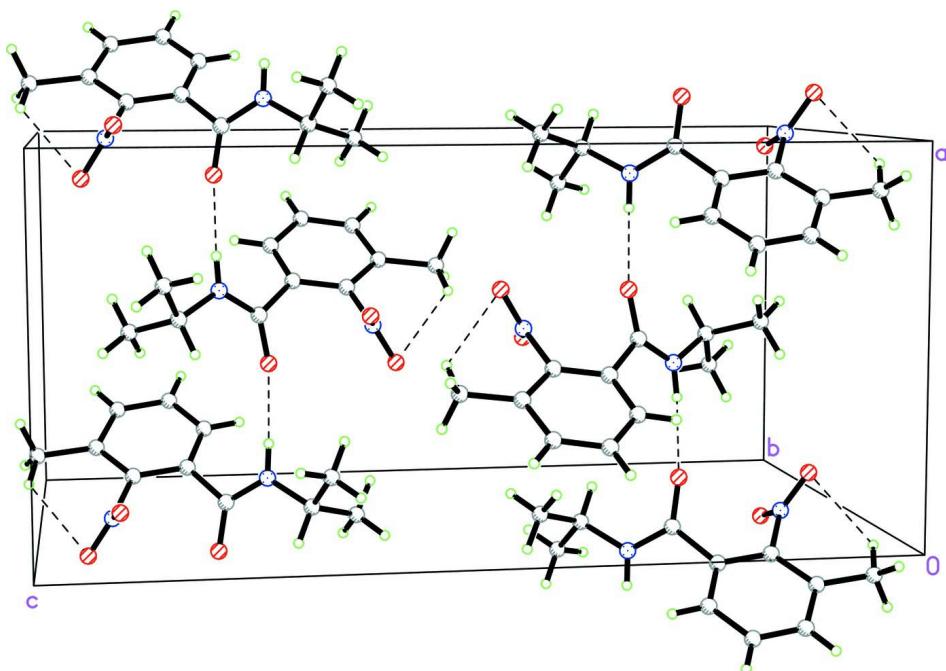
N-isopropyl-3-methyl-2-nitrobenzamide were dissolved in DMF (50 mL). The solution was then poured to ice water. The crystalline product was isolated by filtration, washed with water (600 ml), dried and gave the product 1.8 g. The single crystals were obtained by evaporating the acetone slowly at room temperature for about 14 d.

S3. Refinement

The H atoms were positioned geometrically, with N—H = 0.86 Å (for NH) and C—H = 0.93, 0.96 and 0.98 Å for aromatic, methyl and methine H atoms, respectively, and constrained to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C})$, where $x = 1.5$ for methyl H, and $x = 1.2$ for other H atoms.

**Figure 1**

Asymmetric unit of the title molecular structure, with the atom–labeling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are presented as a small spheres of arbitrary radius.

**Figure 2**

A packing diagram for (I).

N-Isopropyl-3-methyl-2-nitrobenzamide*Crystal data*

$C_{11}H_{14}N_2O_3$
 $M_r = 222.24$
Orthorhombic, $Pbca$
Hall symbol: -P 2ac 2ab
 $a = 9.4230$ (19) Å
 $b = 13.250$ (3) Å
 $c = 20.041$ (4) Å
 $V = 2502.2$ (9) Å³
 $Z = 8$

$F(000) = 944$
 $D_x = 1.180$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 25 reflections
 $\theta = 10\text{--}13^\circ$
 $\mu = 0.09$ mm⁻¹
 $T = 298$ K
Needle, colourless
0.30 × 0.20 × 0.10 mm

Data collection

Enraf–Nonius CAD-4
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator
 $\omega/2\theta$ scans

Absorption correction: ψ scan
(North *et al.*, 1968)

$T_{\min} = 0.974$, $T_{\max} = 0.991$

2260 measured reflections

2260 independent reflections
1135 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.000$
 $\theta_{\max} = 25.3^\circ$, $\theta_{\min} = 2.0^\circ$
 $h = 0 \rightarrow 11$
 $k = 0 \rightarrow 15$
 $l = 0 \rightarrow 24$
3 standard reflections every 200 reflections
intensity decay: 1%

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.066$

$wR(F^2) = 0.174$

$S = 1.00$

2260 reflections

146 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.06P)^2 + 0.9P]$
where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.22$ e Å⁻³

$\Delta\rho_{\min} = -0.16$ e Å⁻³

Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0059 (11)

Special details

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.1002 (2)	0.4262 (2)	0.21752 (13)	0.0801 (9)
N1	-0.0987 (3)	0.4490 (2)	0.27836 (16)	0.0642 (9)

H1A	-0.1881	0.4367	0.2809	0.077*
C1	0.0066 (6)	0.4504 (4)	0.3892 (3)	0.1265 (19)
H1B	0.0699	0.3974	0.3756	0.190*
H1C	-0.0769	0.4215	0.4089	0.190*
H1D	0.0533	0.4928	0.4212	0.190*
O2	-0.0336 (3)	0.51283 (19)	0.09602 (15)	0.0890 (10)
N2	0.0028 (3)	0.4269 (2)	0.08678 (16)	0.0635 (8)
C2	-0.1321 (5)	0.5952 (3)	0.3487 (2)	0.1034 (16)
H2A	-0.1543	0.6345	0.3099	0.155*
H2B	-0.0871	0.6374	0.3814	0.155*
H2C	-0.2178	0.5675	0.3670	0.155*
O3	0.1088 (3)	0.4024 (2)	0.05598 (16)	0.1007 (11)
C3	-0.0342 (4)	0.5115 (3)	0.32994 (19)	0.0676 (11)
H3A	0.0523	0.5415	0.3114	0.081*
C4	-0.0271 (3)	0.4105 (2)	0.22794 (18)	0.0540 (9)
C5	-0.1076 (3)	0.3415 (2)	0.18197 (18)	0.0519 (9)
C6	-0.1956 (4)	0.2662 (3)	0.2058 (2)	0.0698 (11)
H6A	-0.2102	0.2600	0.2516	0.084*
C7	-0.2619 (4)	0.2004 (3)	0.1630 (2)	0.0793 (12)
H7A	-0.3206	0.1500	0.1798	0.095*
C8	-0.2420 (4)	0.2086 (3)	0.0961 (2)	0.0736 (11)
H8A	-0.2884	0.1636	0.0679	0.088*
C9	-0.1545 (4)	0.2819 (3)	0.0684 (2)	0.0642 (10)
C10	-0.0889 (3)	0.3463 (2)	0.11363 (19)	0.0540 (9)
C11	-0.1371 (5)	0.2899 (3)	-0.0058 (2)	0.0973 (15)
H11A	-0.0743	0.3448	-0.0160	0.146*
H11B	-0.2279	0.3020	-0.0260	0.146*
H11C	-0.0981	0.2282	-0.0228	0.146*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0298 (12)	0.130 (2)	0.0802 (19)	-0.0125 (14)	0.0014 (13)	-0.0014 (17)
N1	0.0309 (14)	0.083 (2)	0.079 (2)	-0.0139 (15)	0.0058 (16)	-0.0135 (18)
C1	0.146 (5)	0.126 (4)	0.107 (4)	0.021 (4)	-0.047 (4)	0.009 (4)
O2	0.096 (2)	0.0519 (16)	0.119 (3)	-0.0036 (16)	0.0068 (18)	0.0099 (17)
N2	0.0583 (19)	0.064 (2)	0.068 (2)	-0.0068 (18)	-0.0004 (17)	-0.0036 (17)
C2	0.106 (4)	0.079 (3)	0.125 (4)	0.010 (3)	-0.028 (3)	-0.028 (3)
O3	0.078 (2)	0.114 (2)	0.111 (2)	-0.0197 (18)	0.0431 (19)	-0.0240 (19)
C3	0.053 (2)	0.084 (3)	0.066 (3)	-0.019 (2)	0.001 (2)	-0.005 (2)
C4	0.0306 (17)	0.064 (2)	0.068 (2)	-0.0026 (17)	-0.0016 (18)	0.009 (2)
C5	0.0355 (17)	0.049 (2)	0.071 (2)	-0.0031 (16)	0.0008 (17)	0.0032 (19)
C6	0.058 (2)	0.074 (3)	0.077 (3)	-0.011 (2)	0.002 (2)	0.005 (2)
C7	0.076 (3)	0.065 (3)	0.096 (3)	-0.025 (2)	-0.003 (3)	0.005 (3)
C8	0.070 (3)	0.060 (2)	0.091 (3)	-0.011 (2)	-0.014 (3)	-0.004 (2)
C9	0.061 (2)	0.054 (2)	0.078 (3)	0.0006 (19)	-0.003 (2)	0.000 (2)
C10	0.0430 (19)	0.046 (2)	0.073 (3)	0.0032 (17)	0.0025 (18)	0.0047 (19)
C11	0.115 (4)	0.097 (3)	0.080 (3)	-0.018 (3)	-0.006 (3)	-0.010 (3)

Geometric parameters (\AA , $^{\circ}$)

O1—C4	1.235 (3)	C3—H3A	0.9800
N1—C4	1.318 (4)	C4—C5	1.503 (4)
N1—C3	1.457 (4)	C5—C6	1.382 (4)
N1—H1A	0.8600	C5—C10	1.383 (5)
C1—C3	1.488 (6)	C6—C7	1.375 (5)
C1—H1B	0.9600	C6—H6A	0.9300
C1—H1C	0.9600	C7—C8	1.357 (5)
C1—H1D	0.9600	C7—H7A	0.9300
O2—N2	1.203 (3)	C8—C9	1.391 (5)
N2—O3	1.218 (3)	C8—H8A	0.9300
N2—C10	1.475 (4)	C9—C10	1.391 (5)
C2—C3	1.491 (5)	C9—C11	1.499 (5)
C2—H2A	0.9600	C11—H11A	0.9600
C2—H2B	0.9600	C11—H11B	0.9600
C2—H2C	0.9600	C11—H11C	0.9600
C4—N1—C3	123.4 (3)	N1—C4—C5	116.6 (3)
C4—N1—H1A	118.3	C6—C5—C10	116.9 (3)
C3—N1—H1A	118.3	C6—C5—C4	122.0 (3)
C3—C1—H1B	109.5	C10—C5—C4	120.9 (3)
C3—C1—H1C	109.5	C7—C6—C5	120.9 (4)
H1B—C1—H1C	109.5	C7—C6—H6A	119.5
C3—C1—H1D	109.5	C5—C6—H6A	119.5
H1B—C1—H1D	109.5	C8—C7—C6	120.2 (4)
H1C—C1—H1D	109.5	C8—C7—H7A	119.9
O2—N2—O3	124.3 (3)	C6—C7—H7A	119.9
O2—N2—C10	117.5 (3)	C7—C8—C9	122.2 (4)
O3—N2—C10	118.2 (3)	C7—C8—H8A	118.9
C3—C2—H2A	109.5	C9—C8—H8A	118.9
C3—C2—H2B	109.5	C10—C9—C8	115.6 (4)
H2A—C2—H2B	109.5	C10—C9—C11	123.7 (4)
C3—C2—H2C	109.5	C8—C9—C11	120.7 (4)
H2A—C2—H2C	109.5	C5—C10—C9	124.1 (3)
H2B—C2—H2C	109.5	C5—C10—N2	118.0 (3)
N1—C3—C1	111.4 (3)	C9—C10—N2	117.8 (3)
N1—C3—C2	110.1 (3)	C9—C11—H11A	109.5
C1—C3—C2	111.3 (4)	C9—C11—H11B	109.5
N1—C3—H3A	108.0	H11A—C11—H11B	109.5
C1—C3—H3A	108.0	C9—C11—H11C	109.5
C2—C3—H3A	108.0	H11A—C11—H11C	109.5
O1—C4—N1	124.1 (3)	H11B—C11—H11C	109.5
O1—C4—C5	119.3 (3)	 	
C4—N1—C3—C1	-95.0 (4)	C7—C8—C9—C11	-178.9 (4)
C4—N1—C3—C2	141.1 (4)	C6—C5—C10—C9	1.1 (5)
C3—N1—C4—O1	-3.0 (6)	C4—C5—C10—C9	176.8 (3)

C3—N1—C4—C5	175.6 (3)	C6—C5—C10—N2	179.1 (3)
O1—C4—C5—C6	132.9 (4)	C4—C5—C10—N2	-5.2 (5)
N1—C4—C5—C6	-45.8 (4)	C8—C9—C10—C5	-0.8 (5)
O1—C4—C5—C10	-42.6 (5)	C11—C9—C10—C5	178.0 (4)
N1—C4—C5—C10	138.7 (3)	C8—C9—C10—N2	-178.8 (3)
C10—C5—C6—C7	-0.6 (5)	C11—C9—C10—N2	0.0 (5)
C4—C5—C6—C7	-176.3 (3)	O2—N2—C10—C5	-62.5 (4)
C5—C6—C7—C8	-0.2 (6)	O3—N2—C10—C5	118.6 (4)
C6—C7—C8—C9	0.5 (6)	O2—N2—C10—C9	115.6 (4)
C7—C8—C9—C10	0.0 (6)	O3—N2—C10—C9	-63.3 (4)

Hydrogen-bond geometry (\AA , °)

$D\cdots H$	$D—H$	$H\cdots A$	$D\cdots A$	$D—H\cdots A$
N1—H1A···O1 ⁱ	0.86	2.00	2.855 (3)	173
C11—H11A···O3	0.96	2.37	3.021 (5)	124

Symmetry code: (i) $x-1/2, y, -z+1/2$.