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(E)-Methyl 2-[4-(dimethylamino)benzylidene]hydrazinecarboxylate at 123 K

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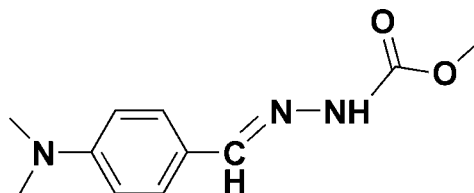
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 Key indicators: single-crystal X-ray study; $T = 123$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.046; wR factor = 0.134; data-to-parameter ratio = 14.6.

The approximately planar molecule of the title compound, $\text{C}_{11}\text{H}_{15}\text{N}_3\text{O}_2$, is in an *E* configuration with respect to the $\text{N}=\text{C}$ double bond. An intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond links the molecules into a one-dimensional chain propagating in the [010] direction.

Related literature

For general background, see: Parashar *et al.* (1988); Hadjoudis *et al.* (1987). For a related structure, see: Shi & Yuan (2006).



Experimental

Crystal data

 $\text{C}_{11}\text{H}_{15}\text{N}_3\text{O}_2$
 $M_r = 221.26$

 Orthorhombic, *Pbca*
 $a = 13.051$ (3) Å

 $b = 9.838$ (2) Å
 $c = 18.637$ (4) Å
 $V = 2392.9$ (9) Å³
 $Z = 8$

 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 123$ (2) K
 $0.29 \times 0.26 \times 0.22$ mm

Data collection

 Bruker SMART CCD diffractometer
 Absorption correction: multi-scan (*SADABS*; Bruker, 2002)
 $T_{\min} = 0.979$, $T_{\max} = 0.981$

 19378 measured reflections
 2111 independent reflections
 1592 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.134$
 $S = 1.11$
 2111 reflections

 145 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.23$ e Å⁻³
 $\Delta\rho_{\min} = -0.27$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N3}-\text{H3}\cdots\text{O1}^i$	0.86	2.16	2.976 (2)	157

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

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAINT* (Bruker, 2002); data reduction: *SAINT*; 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*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB2730).

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supporting information

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(E)-Methyl 2-[4-(dimethylamino)benzylidene]hydrazinecarboxylate at 123 K

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S1. Comment

Benzaldehydehydrazone derivatives have received considerable attention for many years due to their pharmacological activity (Parashar *et al.*, 1988) and their photochromic properties (Hadjoudis *et al.*, 1987). As a further investigation of this type of material, the crystal structure of the title compound, C₁₁H₁₅N₃O₂, (I), is described here.

All the nonhydrogen atoms are coplanar to within ± 0.1429 (14) Å (Fig. 1) in (I). The molecule is in an E conformation with respect to the N=C double bond. The bond lengths and angles of the C=N—N(H)—C groups are similar to those in related compounds (Shi *et al.*, 2006).

An intermolecular N—H \cdots O hydrogen bond (Table 1) links the molecules into a one-dimensional chain aligned along the b direction (Fig. 2).

S2. Experimental

4-(Dimethylamino)benzaldehyde (14.9 g, 0.1 mol) and methyl hydrazinecarboxylate (9.0 g, 0.1 mol) were dissolved in stirred methanol (50 ml) and left for 3 h at room temperature. The resulting solid was filtered off and recrystallized from ethanol to give the title compound in 80% yield. Colourless blocks of (I) were obtained by slow evaporation of a ethanol solution at room temperature (m.p. 452–454 K).

S3. Refinement

The H atoms were geometrically placed (C—H = 0.93–0.96 Å, N—H = 0.86 Å) and refined as riding with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$ or $1.5U_{\text{eq}}(\text{methyl C})$.

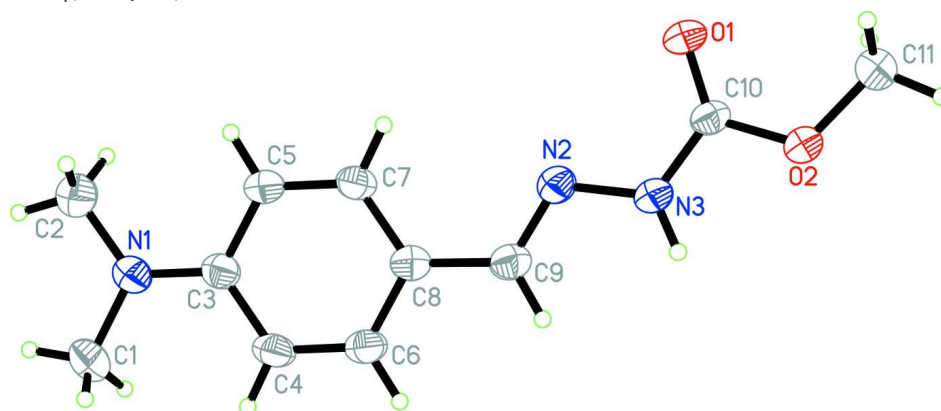
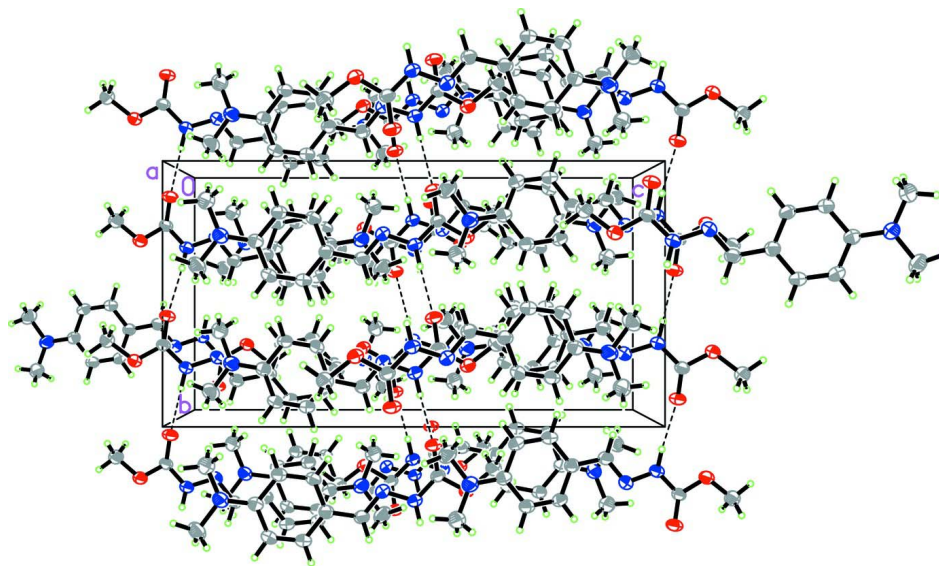


Figure 1

The molecular structure of (I), showing 30% probability displacement ellipsoids for the non-hydrogen atoms.

**Figure 2**

The crystal packing in (II), viewed approximately down the *a* axis with hydrogen bonds indicated by dashed lines.

(E)-Methyl 2-[4-(dimethylamino)benzylidene]hydrazinecarboxylate

Crystal data

$C_{11}H_{15}N_3O_2$

$M_r = 221.26$

Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

$a = 13.051$ (3) Å

$b = 9.838$ (2) Å

$c = 18.637$ (4) Å

$V = 2392.9$ (9) Å³

$Z = 8$

$F(000) = 944$

$D_x = 1.228$ Mg m⁻³

Mo *K*α radiation, $\lambda = 0.71073$ Å

Cell parameters from 2111 reflections

$\theta = 2.2$ – 25.0°

$\mu = 0.09$ mm⁻¹

$T = 123$ K

Block, colourless

$0.29 \times 0.26 \times 0.22$ mm

Data collection

Bruker SMART CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2002)

$T_{\min} = 0.979$, $T_{\max} = 0.981$

19378 measured reflections

2111 independent reflections

1592 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.038$

$\theta_{\max} = 25.0^\circ$, $\theta_{\min} = 2.2^\circ$

$h = -15 \rightarrow 15$

$k = -11 \rightarrow 11$

$l = -21 \rightarrow 22$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.046$

$wR(F^2) = 0.134$

$S = 1.11$

2111 reflections

145 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

$$w = 1/[\sigma^2(F_o^2) + (0.0719P)^2 + 0.3078P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$

$$\Delta\rho_{\max} = 0.23 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.27 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N2	0.27645 (11)	0.23338 (15)	0.56388 (7)	0.0539 (4)
C9	0.29967 (13)	0.13610 (19)	0.60663 (9)	0.0546 (5)
H9	0.2897	0.0469	0.5916	0.066*
O2	0.17803 (10)	0.22366 (12)	0.38955 (6)	0.0610 (4)
N3	0.23964 (12)	0.19224 (14)	0.49757 (7)	0.0554 (4)
H3	0.2363	0.1072	0.4871	0.067*
O1	0.20751 (10)	0.40732 (12)	0.45884 (7)	0.0656 (4)
C3	0.42170 (14)	0.20049 (17)	0.81830 (9)	0.0536 (4)
C8	0.34112 (13)	0.16094 (17)	0.67786 (9)	0.0513 (4)
C5	0.40355 (14)	0.31063 (17)	0.77132 (10)	0.0567 (5)
H5	0.4182	0.3986	0.7865	0.068*
C10	0.20925 (12)	0.28641 (17)	0.44977 (9)	0.0492 (4)
N1	0.45818 (14)	0.22037 (16)	0.88679 (9)	0.0725 (5)
C7	0.36472 (13)	0.29041 (18)	0.70360 (10)	0.0542 (4)
H7	0.3538	0.3652	0.6741	0.065*
C6	0.35985 (14)	0.05251 (18)	0.72400 (10)	0.0593 (5)
H6	0.3453	-0.0352	0.7085	0.071*
C4	0.39921 (15)	0.07084 (18)	0.79189 (10)	0.0621 (5)
H4	0.4111	-0.0046	0.8208	0.075*
C11	0.14121 (16)	0.30910 (19)	0.33264 (10)	0.0667 (5)
H11A	0.1216	0.2538	0.2925	0.100*
H11B	0.1945	0.3707	0.3183	0.100*
H11C	0.0830	0.3599	0.3491	0.100*
C1	0.48056 (19)	0.1055 (2)	0.93250 (11)	0.0833 (7)
H1A	0.5057	0.1373	0.9779	0.125*
H1B	0.5316	0.0494	0.9101	0.125*
H1C	0.4193	0.0534	0.9398	0.125*
C2	0.48463 (18)	0.3522 (2)	0.91387 (11)	0.0787 (6)
H2A	0.5090	0.3439	0.9623	0.118*
H2B	0.4252	0.4098	0.9130	0.118*
H2C	0.5374	0.3912	0.8845	0.118*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N2	0.0641 (9)	0.0466 (9)	0.0509 (9)	0.0017 (6)	-0.0001 (7)	-0.0056 (7)
C9	0.0610 (10)	0.0438 (10)	0.0590 (11)	0.0027 (8)	0.0039 (8)	-0.0028 (8)
O2	0.0838 (9)	0.0425 (7)	0.0568 (8)	0.0004 (6)	-0.0117 (6)	-0.0050 (6)
N3	0.0736 (10)	0.0383 (8)	0.0544 (9)	0.0020 (6)	-0.0048 (7)	-0.0056 (6)
O1	0.0912 (9)	0.0344 (7)	0.0713 (9)	0.0016 (6)	-0.0085 (7)	-0.0070 (6)
C3	0.0545 (10)	0.0476 (10)	0.0588 (11)	0.0041 (7)	-0.0009 (8)	0.0036 (8)
C8	0.0537 (9)	0.0449 (9)	0.0555 (11)	0.0020 (7)	0.0052 (8)	0.0011 (8)
C5	0.0671 (11)	0.0407 (10)	0.0622 (11)	0.0004 (8)	-0.0026 (9)	0.0013 (8)
C10	0.0564 (9)	0.0376 (9)	0.0537 (10)	-0.0007 (7)	0.0042 (8)	-0.0053 (7)
N1	0.0990 (13)	0.0545 (10)	0.0639 (10)	0.0029 (9)	-0.0233 (9)	0.0063 (8)
C7	0.0626 (10)	0.0429 (9)	0.0571 (11)	0.0038 (8)	0.0000 (8)	0.0074 (8)
C6	0.0717 (11)	0.0401 (9)	0.0660 (12)	0.0005 (8)	-0.0003 (9)	0.0013 (8)
C4	0.0780 (12)	0.0444 (10)	0.0639 (12)	0.0062 (9)	-0.0030 (9)	0.0113 (8)
C11	0.0797 (13)	0.0577 (12)	0.0628 (12)	0.0020 (9)	-0.0115 (10)	0.0015 (9)
C1	0.0978 (16)	0.0752 (15)	0.0768 (14)	0.0021 (12)	-0.0256 (12)	0.0202 (11)
C2	0.0936 (15)	0.0716 (14)	0.0708 (13)	-0.0055 (11)	-0.0135 (11)	-0.0009 (12)

Geometric parameters (Å, °)

N2—C9	1.282 (2)	N1—C2	1.434 (3)
N2—N3	1.386 (2)	N1—C1	1.445 (2)
C9—C8	1.454 (2)	C7—H7	0.9300
C9—H9	0.9300	C6—C4	1.377 (3)
O2—C10	1.344 (2)	C6—H6	0.9300
O2—C11	1.436 (2)	C4—H4	0.9300
N3—C10	1.345 (2)	C11—H11A	0.9600
N3—H3	0.8600	C11—H11B	0.9600
O1—C10	1.202 (2)	C11—H11C	0.9600
C3—N1	1.376 (2)	C1—H1A	0.9600
C3—C4	1.398 (2)	C1—H1B	0.9600
C3—C5	1.413 (2)	C1—H1C	0.9600
C8—C6	1.392 (2)	C2—H2A	0.9600
C8—C7	1.395 (2)	C2—H2B	0.9600
C5—C7	1.374 (3)	C2—H2C	0.9600
C5—H5	0.9300		
C9—N2—N3	114.71 (15)	C8—C7—H7	119.0
N2—C9—C8	122.00 (16)	C4—C6—C8	122.19 (17)
N2—C9—H9	119.0	C4—C6—H6	118.9
C8—C9—H9	119.0	C8—C6—H6	118.9
C10—O2—C11	116.70 (13)	C6—C4—C3	121.41 (16)
C10—N3—N2	119.44 (14)	C6—C4—H4	119.3
C10—N3—H3	120.3	C3—C4—H4	119.3
N2—N3—H3	120.3	O2—C11—H11A	109.5
N1—C3—C4	121.92 (16)	O2—C11—H11B	109.5

N1—C3—C5	121.59 (16)	H11A—C11—H11B	109.5
C4—C3—C5	116.49 (16)	O2—C11—H11C	109.5
C6—C8—C7	116.64 (16)	H11A—C11—H11C	109.5
C6—C8—C9	120.04 (16)	H11B—C11—H11C	109.5
C7—C8—C9	123.32 (16)	N1—C1—H1A	109.5
C7—C5—C3	121.32 (16)	N1—C1—H1B	109.5
C7—C5—H5	119.3	H1A—C1—H1B	109.5
C3—C5—H5	119.3	N1—C1—H1C	109.5
O1—C10—O2	124.50 (16)	H1A—C1—H1C	109.5
O1—C10—N3	126.45 (16)	H1B—C1—H1C	109.5
O2—C10—N3	109.03 (14)	N1—C2—H2A	109.5
C3—N1—C2	122.56 (16)	N1—C2—H2B	109.5
C3—N1—C1	120.36 (17)	H2A—C2—H2B	109.5
C2—N1—C1	116.83 (17)	N1—C2—H2C	109.5
C5—C7—C8	121.95 (16)	H2A—C2—H2C	109.5
C5—C7—H7	119.0	H2B—C2—H2C	109.5

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N3—H3...O1 ⁱ	0.86	2.16	2.976 (2)	157

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