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(E)-Methyl N'-[1-(4-methylphenyl)ethylidene]hydrazinecarboxylate

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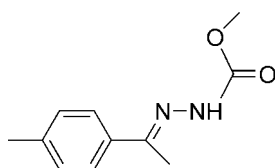
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 Key indicators: single-crystal X-ray study; $T = 273$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.057; wR factor = 0.172; data-to-parameter ratio = 14.1.

The title molecule, $\text{C}_{11}\text{H}_{14}\text{N}_2\text{O}_2$, adopts a *trans* configuration with respect to the $\text{C}=\text{N}$ bond. The dihedral angle between the benzene ring and the hydrazinecarboxylate plane is $7.61(16)^\circ$. In the crystal structure, molecules are linked into centrosymmetric dimers by $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds and the dimers are linked together by $\text{C}-\text{H}\cdots\pi$ interactions.

Related literature

For general background, see: Parashar *et al.* (1988); Hadjoudis *et al.* (1987); Borg *et al.* (1999). For a related structure, see Lv *et al.* (2008).



Experimental

Crystal data

 $\text{C}_{11}\text{H}_{14}\text{N}_2\text{O}_2$
 $M_r = 206.24$
 Monoclinic, $P2_1/c$
 $a = 11.5197(3)$ Å
 $b = 5.5734(6)$ Å
 $c = 17.3281(2)$ Å

 $\beta = 94.193(14)^\circ$
 $V = 1109.55(12)$ Å³
 $Z = 4$
 Mo $K\alpha$ radiation

 $\mu = 0.09$ mm⁻¹
 $T = 273(2)$ K
 $0.24 \times 0.22 \times 0.20$ mm

Data collection

 Bruker SMART CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2002)
 $T_{\min} = 0.978$, $T_{\max} = 0.980$
 5502 measured reflections
 1951 independent reflections
 1209 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.057$
 $wR(F^2) = 0.172$
 $S = 1.03$
 1951 reflections
 138 parameters

 3 restraints
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.28$ e Å⁻³
 $\Delta\rho_{\min} = -0.26$ e Å⁻³
Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N2}-\text{H2A}\cdots\text{O1}^{\text{i}}$	0.86	2.12	2.944 (3)	162
$\text{C2}-\text{H2}\cdots\text{Cg1}^{\text{ii}}$	0.93	2.83	3.538 (3)	134

 Symmetry codes: (i) $-x, -y, -z + 1$; (ii) $-x + 1, y + \frac{1}{2}, -z + \frac{1}{2}$. Cg1 is the centroid of the C2–C7 benzene ring.

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: CV2459).

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supporting information

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(E)-Methyl N'-[1-(4-methylphenyl)ethylidene]hydrazinecarboxylate

Lu-Ping Lv, Wen-Bo Yu, Feng Wang, Wei-Wei Li and Xian-Chao Hu

S1. Comment

Benzaldehydhydrazone derivatives have received considerable attention for a long time, due to their pharmacological activities (Parashar *et al.*, 1988) and their photochromic properties (Hadjoudis *et al.*, 1987). They are important intermediates for 1,3,4-oxadiazoles, which have been reported to be versatile compounds with many useful properties (Borg *et al.*, 1999). As a further investigation of this type of derivatives, we report herein the crystal structure of the title compound.

The title molecule (Fig. 1) adopts a trans configuration with respect to the C=N double bond. The bond lengths and angles are comparable to those observed for (E)-methyl N'-[1-(4-methoxyphenyl)ethylidene]hydrazinecarboxylate (Lv *et al.*, 2008). All atoms of O1/O2/N1/N2/C8-C11 are coplanar within ± 0.093 (2) Å. The dihedral angle between the benzene (C2-C7) and O1/O2/N1/N2/C8-C11 planes is 7.61 (16)°.

In the crystal structure, intermolecular N—H \cdots O hydrogen bonds (Table 1) link the molecules into centrosymmetric dimers (Fig. 2). A C—H \cdots π contact (Table 1) between the benzene ring (centroid Cg1) and H atom of aromatic C2 further stabilizes the structure.

S2. Experimental

4-Methyl-acetophenone (1.34 g, 0.01 mol) and methyl hydrazinecarboxylate (0.90 g, 0.01 mol) were dissolved in stirred methanol (25 ml) and left for 5.5 h at room temperature. The resulting solid was filtered off and recrystallized from ethanol to give the title compound (yield 93%, m.p. 453–455 K). Single crystals of the title compound suitable for X-ray analysis were obtained by slow evaporation of an ethanol solution.

S3. Refinement

H atoms were positioned geometrically, with N-H = 0.86 Å and C-H = 0.93 and 0.96 Å for aromatic and methyl H, respectively, and constrained to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C}, \text{N})$, where $x = 1.5$ for methyl H and $x = 1.2$ for all other H atoms. A rotating group model was used for the methyl groups.

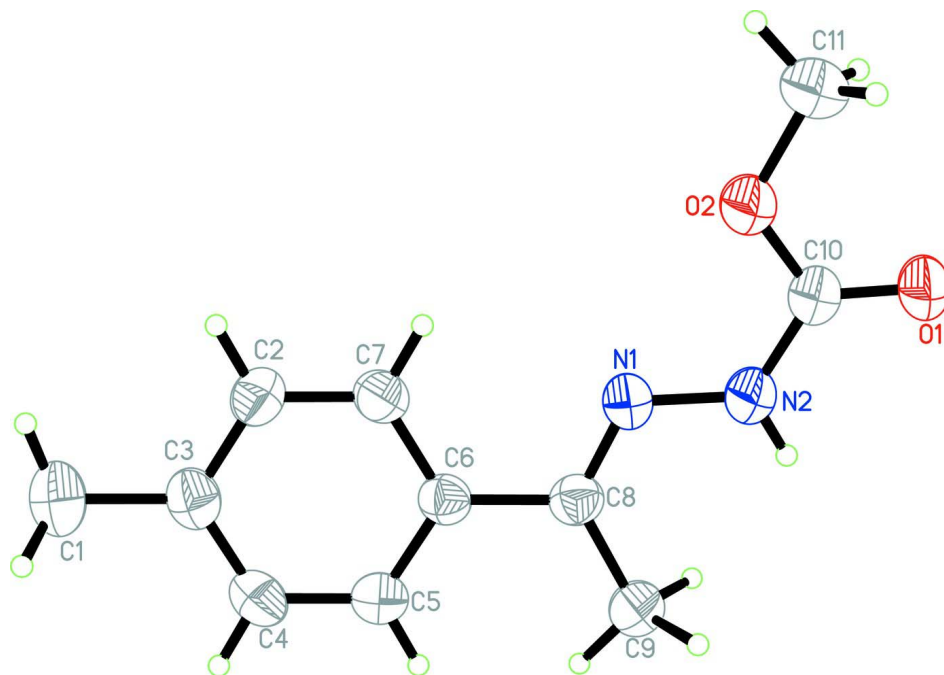


Figure 1

The molecular structure of the title compound, with the atom-numbering scheme. Displacement ellipsoids are drawn at the 40% probability level.

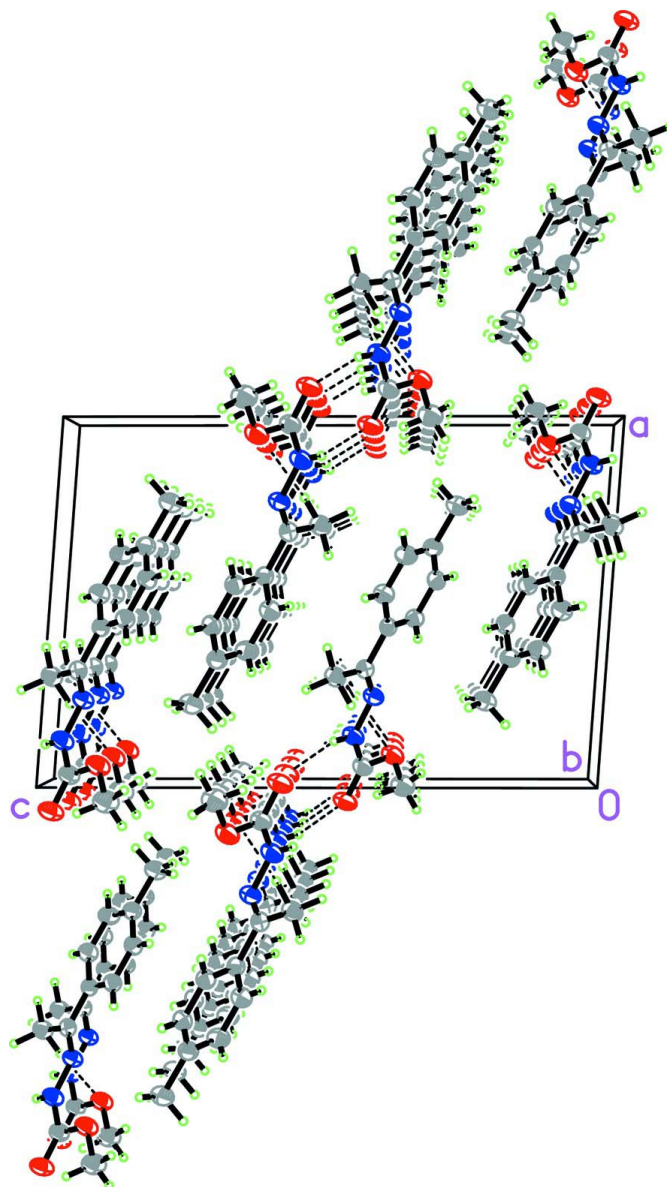


Figure 2

A packing diagram of the title compound. Hydrogen bonds are shown as dashed lines.

(E)-Methyl N'-[1-(4-methylphenyl)ethylidene]hydrazinecarboxylate

Crystal data

$C_{11}H_{14}N_2O_2$

$M_r = 206.24$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2_1/c$

$a = 11.5197(3)\ \text{\AA}$

$b = 5.5734(6)\ \text{\AA}$

$c = 17.3281(2)\ \text{\AA}$

$\beta = 94.193(14)^\circ$

$V = 1109.55(12)\ \text{\AA}^3$

$Z = 4$

$F(000) = 440$

$D_x = 1.235\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 1951 reflections

$\theta = 1.8\text{--}25.0^\circ$

$\mu = 0.09\ \text{mm}^{-1}$

$T = 273\ \text{K}$

Block, colourless

$0.24 \times 0.22 \times 0.20\ \text{mm}$

Data collection

Bruker SMART CCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 φ and ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2002)
 $T_{\min} = 0.978$, $T_{\max} = 0.980$

5502 measured reflections
1951 independent reflections
1209 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$
 $\theta_{\text{max}} = 25.0^\circ$, $\theta_{\text{min}} = 1.8^\circ$
 $h = -13 \rightarrow 13$
 $k = -6 \rightarrow 6$
 $l = -19 \rightarrow 20$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.057$
 $wR(F^2) = 0.172$
 $S = 1.03$
1951 reflections
138 parameters
3 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
 $w = 1/[\sigma^2(F_o^2) + (0.07P)^2 + 0.5722P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} < 0.001$
 $\Delta\rho_{\text{max}} = 0.28 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.26 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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 $F^2 > 2\sigma(F^2)$ 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>	$U_{\text{iso}}^*/U_{\text{eq}}$
C6	0.4332 (2)	-0.0332 (5)	0.39288 (14)	0.0447 (6)
C5	0.5218 (2)	-0.2016 (5)	0.40434 (17)	0.0571 (8)
H5	0.5107	-0.3340	0.4357	0.069*
C8	0.3200 (2)	-0.0595 (5)	0.42808 (15)	0.0472 (7)
C7	0.4558 (2)	0.1643 (5)	0.34621 (16)	0.0520 (7)
H7	0.3994	0.2827	0.3379	0.062*
C3	0.6471 (2)	0.0163 (5)	0.32361 (16)	0.0548 (7)
C2	0.5594 (2)	0.1865 (5)	0.31268 (16)	0.0548 (7)
H2	0.5712	0.3193	0.2817	0.066*
C10	0.0517 (2)	0.2495 (5)	0.42066 (17)	0.0562 (7)
C4	0.6258 (2)	-0.1771 (5)	0.37036 (18)	0.0632 (8)
H4	0.6830	-0.2937	0.3792	0.076*
C1	0.7601 (3)	0.0414 (7)	0.28540 (19)	0.0764 (10)
H1A	0.7559	0.1786	0.2518	0.115*
H1B	0.7732	-0.1002	0.2557	0.115*
H1C	0.8230	0.0618	0.3243	0.115*

C9	0.3016 (2)	-0.2711 (5)	0.47910 (17)	0.0602 (8)
H9A	0.2698	-0.2178	0.5258	0.063*
H9C	0.3746	-0.3499	0.4916	0.063*
H9B	0.2484	-0.3812	0.4526	0.063*
C11	-0.0088 (3)	0.5827 (6)	0.3454 (2)	0.0789 (10)
H11A	-0.0725	0.5048	0.3167	0.118*
H11B	0.0235	0.7033	0.3137	0.118*
H11C	-0.0362	0.6564	0.3907	0.118*
N2	0.13765 (19)	0.0920 (4)	0.44132 (14)	0.0613 (7)
H2A	0.1257	-0.0182	0.4746	0.074*
N1	0.24476 (18)	0.1054 (4)	0.40979 (13)	0.0548 (6)
O2	0.07955 (17)	0.4080 (4)	0.36804 (12)	0.0690 (7)
O1	-0.04293 (16)	0.2424 (4)	0.44769 (13)	0.0733 (7)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C6	0.0480 (15)	0.0439 (15)	0.0421 (14)	0.0020 (12)	0.0018 (11)	-0.0030 (12)
C5	0.0586 (17)	0.0476 (17)	0.0660 (19)	0.0056 (13)	0.0102 (14)	0.0060 (14)
C8	0.0514 (15)	0.0426 (15)	0.0478 (15)	0.0001 (13)	0.0055 (12)	0.0006 (12)
C7	0.0509 (16)	0.0495 (17)	0.0559 (17)	0.0076 (13)	0.0054 (13)	0.0025 (14)
C3	0.0485 (16)	0.0588 (18)	0.0572 (18)	-0.0005 (14)	0.0052 (13)	-0.0077 (15)
C2	0.0564 (17)	0.0573 (18)	0.0515 (17)	-0.0049 (14)	0.0089 (13)	0.0058 (14)
C10	0.0485 (16)	0.0579 (18)	0.0631 (19)	-0.0015 (14)	0.0098 (14)	0.0062 (15)
C4	0.0537 (17)	0.0569 (19)	0.080 (2)	0.0145 (14)	0.0090 (15)	0.0013 (16)
C1	0.0543 (18)	0.095 (3)	0.082 (2)	-0.0016 (18)	0.0152 (16)	-0.004 (2)
C9	0.0566 (17)	0.0590 (19)	0.0662 (19)	0.0039 (14)	0.0116 (14)	0.0098 (15)
C11	0.071 (2)	0.070 (2)	0.096 (3)	0.0151 (18)	0.0100 (18)	0.0236 (19)
N2	0.0532 (14)	0.0602 (16)	0.0729 (16)	0.0043 (12)	0.0202 (12)	0.0171 (13)
N1	0.0459 (13)	0.0557 (15)	0.0641 (15)	0.0030 (11)	0.0133 (11)	0.0075 (12)
O2	0.0572 (12)	0.0718 (15)	0.0801 (15)	0.0104 (11)	0.0194 (11)	0.0241 (12)
O1	0.0492 (12)	0.0806 (16)	0.0922 (16)	0.0046 (11)	0.0205 (11)	0.0214 (13)

Geometric parameters (Å, °)

C6—C5	1.390 (4)	C10—N2	1.351 (4)
C6—C7	1.402 (4)	C4—H4	0.9300
C6—C8	1.487 (3)	C1—H1A	0.9600
C5—C4	1.380 (4)	C1—H1B	0.9600
C5—H5	0.9300	C1—H1C	0.9600
C8—N1	1.286 (3)	C9—H9A	0.9600
C8—C9	1.498 (4)	C9—H9C	0.9600
C7—C2	1.370 (4)	C9—H9B	0.9600
C7—H7	0.9300	C11—O2	1.442 (3)
C3—C4	1.381 (4)	C11—H11A	0.9600
C3—C2	1.389 (4)	C11—H11B	0.9600
C3—C1	1.509 (4)	C11—H11C	0.9600
C2—H2	0.9300	N2—N1	1.388 (3)

C10—O1	1.218 (3)	N2—H2A	0.8600
C10—O2	1.326 (3)		
C5—C6—C7	116.5 (2)	C3—C1—H1A	109.5
C5—C6—C8	122.1 (2)	C3—C1—H1B	109.5
C7—C6—C8	121.3 (2)	H1A—C1—H1B	109.5
C4—C5—C6	121.6 (3)	C3—C1—H1C	109.5
C4—C5—H5	119.2	H1A—C1—H1C	109.5
C6—C5—H5	119.2	H1B—C1—H1C	109.5
N1—C8—C6	115.1 (2)	C8—C9—H9A	109.5
N1—C8—C9	125.8 (2)	C8—C9—H9C	109.5
C6—C8—C9	119.1 (2)	H9A—C9—H9C	109.5
C2—C7—C6	121.3 (3)	C8—C9—H9B	109.5
C2—C7—H7	119.3	H9A—C9—H9B	109.5
C6—C7—H7	119.3	H9C—C9—H9B	109.5
C4—C3—C2	117.0 (3)	O2—C11—H11A	109.5
C4—C3—C1	121.7 (3)	O2—C11—H11B	109.5
C2—C3—C1	121.3 (3)	H11A—C11—H11B	109.5
C7—C2—C3	121.9 (3)	O2—C11—H11C	109.5
C7—C2—H2	119.1	H11A—C11—H11C	109.5
C3—C2—H2	119.1	H11B—C11—H11C	109.5
O1—C10—O2	123.7 (3)	C10—N2—N1	121.1 (2)
O1—C10—N2	122.4 (3)	C10—N2—H2A	119.5
O2—C10—N2	113.9 (2)	N1—N2—H2A	119.5
C5—C4—C3	121.6 (3)	C8—N1—N2	117.8 (2)
C5—C4—H4	119.2	C10—O2—C11	115.8 (2)
C3—C4—H4	119.2		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N2—H2A \cdots O1 ⁱ	0.86	2.12	2.944 (3)	162
C2—H2 \cdots Cg1 ⁱⁱ	0.93	2.83	3.538 (3)	134

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