

(*E*)-1-[4-(Dimethylamino)benzylidene]-semicarbazide–acetic acid (1/2)

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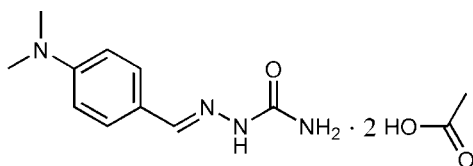
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Key indicators: single-crystal X-ray study; $T = 294$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.062; wR factor = 0.202; data-to-parameter ratio = 13.6.

In the crystal structure of the title compound, $\text{C}_{10}\text{H}_{16}\text{N}_4\text{O} \cdot 2\text{C}_2\text{H}_4\text{O}_2$, the molecules interact by way of $\text{O}-\text{H} \cdots \text{O}$ and $\text{N}-\text{H} \cdots \text{O}$ hydrogen bonds, resulting in [010] chains built up from the 1-[4-(dimethylamino)benzylidene]semicarbazide molecule and one of the acetic acid molecules. The other acetic acid molecule forms isolated centrosymmetric dimers.

Related literature

For a related structure, see Tai *et al.* (2007).



Experimental

Crystal data

$\text{C}_{10}\text{H}_{14}\text{N}_4\text{O} \cdot 2\text{C}_2\text{H}_4\text{O}_2$
 $M_r = 326.36$
 Triclinic, $P\bar{1}$
 $a = 7.2931$ (2) Å
 $b = 10.7952$ (8) Å
 $c = 12.2472$ (7) Å

$\alpha = 71.422$ (17)°
 $\beta = 80.020$ (4)°
 $\gamma = 73.947$ (4)°
 $V = 874.49$ (12) Å³
 $Z = 2$
 Mo $K\alpha$ radiation

$\mu = 0.10$ mm⁻¹
 $T = 294$ (2) K

0.16 × 0.14 × 0.12 mm

Data collection

Bruker SMART CCD diffractometer
 Absorption correction: multi-scan (*SADABS*; Bruker, 1997)
 $T_{\min} = 0.985$, $T_{\max} = 0.989$

4441 measured reflections
 3064 independent reflections
 1351 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.040$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.063$
 $wR(F^2) = 0.202$
 $S = 1.01$
 3064 reflections
 225 parameters
 1 restraint

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.19$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

<i>D</i> — <i>H</i> ··· <i>A</i>	<i>D</i> — <i>H</i>	<i>H</i> ··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> — <i>H</i> ··· <i>A</i>
O3—H3···O2 ⁱ	0.85	1.82	2.658 (6)	170
O5—H5A···O1 ⁱⁱ	0.82	1.78	2.579 (4)	166
N4—H4A···O4 ⁱⁱⁱ	0.87 (4)	2.48 (4)	3.164 (6)	135 (3)
N3—H3A···O1 ^{iv}	0.897 (10)	2.017 (11)	2.913 (5)	178 (4)
N4—H4B···O4 ^v	0.91 (6)	2.16 (6)	3.051 (6)	166 (5)

Symmetry codes: (i) $-x + 1, -y + 1, -z + 1$; (ii) $x + 1, y, z$; (iii) $-x + 1, -y + 2, -z + 1$; (iv) $-x, -y + 1, -z + 1$; (v) $x - 1, y, z$.

Data collection: *SMART* (Bruker, 1997); cell refinement: *SAINTE* (Bruker, 1997); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1997); 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: HB2411).

References

- Bruker (1997). *SADABS* (Version 2.01), *SMART* (Version 5.044), *SAINTE* (Version 5.01) and *SHELXTL* (Version 5.10). Bruker AXS Inc., Madison, Wisconsin, USA.
 Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.
 Tai, X.-S., Hao, M.-Y., Yin, J. & Liang, Z.-P. (2007). *Acta Cryst.* **E63**, o1725–o1726.

supplementary materials

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(*E*)-1-[4-(Dimethylamino)benzylidene]semicarbazide-acetic acid (1/2)

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Comment

Their preparation, properties and applications of Schiff bases are important in the development of coordination chemistry. The asymmetric unit of the title compound, (I), contains one (*E*)-1-(4-(dimethylamino)benzylidene)semicarbazide molecule and two acetic acid molecules (Fig. 1). The bond lengths and angles of the Schiff base in (I) agree with those in the related (*E*)-1-(4-hydroxybenzylidene)semicarbazide hemihydrate (Tai *et al.*, 2007). The main molecule in (I) is essentially planar, with a maximum deviation from the mean plane for the non-hydrogen atoms of 0.042 (2) Å. The crystal structure of (I) is stabilized by O—H...O and N—H...O hydrogen bonds (Fig. 2 and Table 1), to result in chains built up from the C₁₀H₁₆N₄O molecule and the C13-containing acetic acid molecule. Conversely, the C11 acetic acid molecule forms isolated inversion dimers.

Experimental

A mixture of 4-(dimethylamino)benzaldehyde (0.01 mol) and semicarbazide hydrochloride (0.01 mol) in ethanol (10 ml) was refluxed for 1 h. After cooling, filtration and drying, the compound (*E*)-1-(4-(dimethylamino)benzylidene)semicarbazide was obtained. 10 mg of this compound was dissolved in acetic acid (8 ml), and the solution was then allowed to evaporate at room temperature; light yellow blocks of (I) were formed after 12 d.

Refinement

The N-bound H atoms were located in a difference map and their positions and U_{iso} values were freely refined.

The O-bound H atoms were located in a difference map and refined as riding in their as-found relative positions with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$.

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

Figures

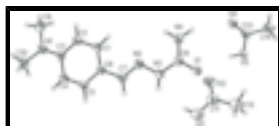


Fig. 1. The molecular structure of (I), drawn with 30% probability ellipsoids (arbitrary spheres for the H atoms).

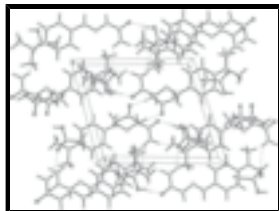


Fig. 2. The crystal packing of (I), viewed along *a* axis. Hydrogen bonds are indicated by dashed lines.

(*E*)-1-[4-(Dimethylamino)benzylidene]semicarbazide–acetic acid (1/2)

Crystal data

$C_{10}H_{14}N_4O \cdot 2C_2H_4O_2$	$Z = 2$
$M_r = 326.36$	$F_{000} = 348$
Triclinic, <i>PT</i>	$D_x = 1.239 \text{ Mg m}^{-3}$
Hall symbol: -P 1	Mo $K\alpha$ radiation
$a = 7.2931 (2) \text{ \AA}$	$\lambda = 0.71073 \text{ \AA}$
$b = 10.7952 (8) \text{ \AA}$	Cell parameters from 886 reflections
$c = 12.2472 (7) \text{ \AA}$	$\theta = 2.9\text{--}23.5^\circ$
$\alpha = 71.422 (17)^\circ$	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 80.020 (4)^\circ$	$T = 294 (2) \text{ K}$
$\gamma = 73.947 (4)^\circ$	Block, light yellow
$V = 874.49 (12) \text{ \AA}^3$	$0.16 \times 0.14 \times 0.12 \text{ mm}$

Data collection

Bruker SMART CCD diffractometer	3064 independent reflections
Radiation source: fine-focus sealed tube	1351 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.040$
$T = 294(2) \text{ K}$	$\theta_{\text{max}} = 25.0^\circ$
φ and ω scans	$\theta_{\text{min}} = 1.8^\circ$
Absorption correction: multi-scan (SADABS; Bruker, 1997)	$h = -8 \rightarrow 7$
$T_{\text{min}} = 0.985$, $T_{\text{max}} = 0.989$	$k = -12 \rightarrow 12$
4441 measured reflections	$l = -14 \rightarrow 6$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.063$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.202$	$w = 1/[\sigma^2(F_o^2) + (0.0848P)^2 + 0.1483P]$
$S = 1.01$	where $P = (F_o^2 + 2F_c^2)/3$
	$(\Delta/\sigma)_{\text{max}} = 0.005$

3064 reflections $\Delta\rho_{\max} = 0.19 \text{ e \AA}^{-3}$
 225 parameters $\Delta\rho_{\min} = -0.21 \text{ e \AA}^{-3}$
 1 restraint Extinction correction: none
 Primary atom site location: structure-invariant direct methods

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 $F^2 > 2\sigma(F^2)$ 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>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	-0.0214 (4)	0.6237 (2)	0.5764 (2)	0.0645 (9)
N1	0.3989 (6)	0.9089 (3)	-0.2633 (3)	0.0756 (12)
N2	0.1380 (5)	0.7164 (3)	0.2810 (3)	0.0543 (9)
N3	0.0888 (5)	0.6399 (3)	0.3916 (3)	0.0609 (10)
N4	0.0405 (6)	0.8213 (4)	0.4616 (4)	0.0686 (12)
C1	0.2324 (6)	0.8560 (4)	0.0449 (3)	0.0593 (12)
H1	0.1941	0.9067	0.0972	0.071*
C2	0.2830 (6)	0.9184 (4)	-0.0688 (3)	0.0621 (13)
H2	0.2769	1.0101	-0.0915	0.074*
C3	0.3434 (6)	0.8470 (4)	-0.1512 (3)	0.0540 (11)
C4	0.3461 (6)	0.7089 (4)	-0.1113 (3)	0.0549 (12)
H4	0.3838	0.6576	-0.1631	0.066*
C5	0.2941 (6)	0.6479 (3)	0.0034 (3)	0.0526 (11)
H5	0.2977	0.5565	0.0268	0.063*
C6	0.2366 (5)	0.7196 (3)	0.0842 (3)	0.0470 (10)
C7	0.1849 (6)	0.6532 (4)	0.2048 (3)	0.0537 (11)
H7	0.1862	0.5622	0.2267	0.064*
C8	0.0324 (6)	0.6951 (4)	0.4796 (3)	0.0520 (11)
C9	0.3996 (8)	1.0509 (4)	-0.3013 (4)	0.0873 (17)
H9A	0.2720	1.1034	-0.2895	0.131*
H9B	0.4438	1.0761	-0.3820	0.131*
H9C	0.4833	1.0670	-0.2575	0.131*
C10	0.4361 (8)	0.8414 (5)	-0.3535 (4)	0.0910 (17)
H10A	0.5516	0.7714	-0.3415	0.137*
H10B	0.4506	0.9053	-0.4278	0.137*
H10C	0.3308	0.8030	-0.3508	0.137*
O2	0.4506 (5)	0.5408 (3)	0.6268 (3)	0.0902 (12)

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O3	0.4009 (5)	0.3657 (3)	0.5886 (3)	0.0960 (12)
H3	0.4495	0.4040	0.5221	0.115*
C11	0.3914 (7)	0.4363 (5)	0.6571 (5)	0.0765 (14)
C12	0.3084 (8)	0.3842 (5)	0.7785 (5)	0.0920 (16)
H12A	0.4089	0.3254	0.8251	0.138*
H12B	0.2162	0.3355	0.7789	0.138*
H12C	0.2468	0.4583	0.8096	0.138*
O4	0.9048 (5)	0.8908 (3)	0.6896 (3)	0.0806 (10)
O5	0.8966 (5)	0.6762 (3)	0.7728 (2)	0.0788 (10)
H5A	0.9164	0.6733	0.7056	0.118*
C13	0.8785 (6)	0.8016 (4)	0.7749 (4)	0.0606 (12)
C14	0.8229 (7)	0.8142 (5)	0.8945 (4)	0.0776 (15)
H14A	0.9334	0.8155	0.9264	0.116*
H14B	0.7724	0.7390	0.9418	0.116*
H14C	0.7270	0.8961	0.8924	0.116*
H4A	0.070 (5)	0.872 (3)	0.393 (3)	0.046 (11)*
H3A	0.070 (5)	0.558 (2)	0.400 (3)	0.070 (13)*
H4B	0.000 (8)	0.857 (6)	0.523 (5)	0.14 (2)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.102 (2)	0.0522 (15)	0.0443 (17)	-0.0302 (15)	0.0101 (16)	-0.0185 (13)
N1	0.121 (4)	0.064 (2)	0.047 (2)	-0.036 (2)	0.012 (2)	-0.0201 (18)
N2	0.068 (2)	0.0523 (18)	0.047 (2)	-0.0230 (17)	0.0054 (18)	-0.0180 (17)
N3	0.091 (3)	0.0506 (19)	0.047 (2)	-0.0283 (19)	0.0085 (19)	-0.0198 (17)
N4	0.109 (4)	0.054 (2)	0.046 (2)	-0.033 (2)	0.002 (2)	-0.012 (2)
C1	0.080 (3)	0.047 (2)	0.053 (3)	-0.014 (2)	0.005 (2)	-0.023 (2)
C2	0.089 (4)	0.044 (2)	0.056 (3)	-0.022 (2)	0.009 (3)	-0.019 (2)
C3	0.066 (3)	0.054 (2)	0.044 (2)	-0.018 (2)	0.002 (2)	-0.0173 (19)
C4	0.073 (3)	0.047 (2)	0.049 (2)	-0.015 (2)	0.003 (2)	-0.0227 (19)
C5	0.066 (3)	0.0389 (19)	0.053 (3)	-0.0144 (19)	-0.001 (2)	-0.0133 (18)
C6	0.051 (3)	0.047 (2)	0.043 (2)	-0.0113 (19)	0.000 (2)	-0.0166 (18)
C7	0.061 (3)	0.046 (2)	0.056 (3)	-0.017 (2)	0.001 (2)	-0.017 (2)
C8	0.066 (3)	0.045 (2)	0.048 (3)	-0.018 (2)	-0.002 (2)	-0.014 (2)
C9	0.129 (5)	0.066 (3)	0.057 (3)	-0.032 (3)	0.011 (3)	-0.006 (2)
C10	0.135 (5)	0.102 (4)	0.049 (3)	-0.047 (3)	0.013 (3)	-0.035 (3)
O2	0.106 (3)	0.067 (2)	0.113 (3)	-0.038 (2)	-0.008 (2)	-0.0342 (19)
O3	0.125 (3)	0.074 (2)	0.104 (3)	-0.041 (2)	-0.011 (2)	-0.030 (2)
C11	0.067 (4)	0.068 (3)	0.098 (4)	-0.010 (3)	-0.020 (3)	-0.026 (3)
C12	0.085 (4)	0.095 (4)	0.101 (4)	-0.026 (3)	-0.011 (3)	-0.031 (3)
O4	0.116 (3)	0.0558 (18)	0.075 (2)	-0.0313 (18)	0.005 (2)	-0.0225 (16)
O5	0.130 (3)	0.0568 (17)	0.0556 (19)	-0.0280 (17)	0.006 (2)	-0.0264 (15)
C13	0.067 (3)	0.055 (2)	0.067 (3)	-0.017 (2)	-0.002 (3)	-0.027 (2)
C14	0.095 (4)	0.085 (3)	0.072 (3)	-0.029 (3)	0.002 (3)	-0.047 (3)

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

O1—C8	1.256 (4)	C7—H7	0.9300
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N1—C3	1.367 (5)	C9—H9A	0.9600
N1—C9	1.454 (5)	C9—H9B	0.9600
N1—C10	1.458 (5)	C9—H9C	0.9600
N2—C7	1.273 (4)	C10—H10A	0.9600
N2—N3	1.388 (4)	C10—H10B	0.9600
N3—C8	1.344 (5)	C10—H10C	0.9600
N3—H3A	0.897 (10)	O2—C11	1.243 (5)
N4—C8	1.325 (5)	O3—C11	1.284 (5)
N4—H4A	0.87 (4)	O3—H3	0.8522
N4—H4B	0.91 (6)	C11—C12	1.496 (7)
C1—C2	1.375 (5)	C12—H12A	0.9600
C1—C6	1.389 (5)	C12—H12B	0.9600
C1—H1	0.9300	C12—H12C	0.9600
C2—C3	1.401 (5)	O4—C13	1.203 (5)
C2—H2	0.9300	O5—C13	1.331 (5)
C3—C4	1.409 (5)	O5—H5A	0.8200
C4—C5	1.384 (5)	C13—C14	1.490 (6)
C4—H4	0.9300	C14—H14A	0.9600
C5—C6	1.386 (5)	C14—H14B	0.9600
C5—H5	0.9300	C14—H14C	0.9600
C6—C7	1.455 (5)		
?...?	?		
C3—N1—C9	121.4 (3)	N4—C8—N3	118.8 (4)
C3—N1—C10	121.8 (4)	N1—C9—H9A	109.5
C9—N1—C10	116.4 (4)	N1—C9—H9B	109.5
C7—N2—N3	114.9 (3)	H9A—C9—H9B	109.5
C8—N3—N2	120.9 (3)	N1—C9—H9C	109.5
C8—N3—H3A	119 (3)	H9A—C9—H9C	109.5
N2—N3—H3A	119 (3)	H9B—C9—H9C	109.5
C8—N4—H4A	122 (2)	N1—C10—H10A	109.5
C8—N4—H4B	117 (4)	N1—C10—H10B	109.5
H4A—N4—H4B	121 (4)	H10A—C10—H10B	109.5
C2—C1—C6	122.1 (3)	N1—C10—H10C	109.5
C2—C1—H1	118.9	H10A—C10—H10C	109.5
C6—C1—H1	118.9	H10B—C10—H10C	109.5
C1—C2—C3	121.6 (4)	C11—O3—H3	108.8
C1—C2—H2	119.2	O2—C11—O3	122.9 (5)
C3—C2—H2	119.2	O2—C11—C12	120.9 (5)
N1—C3—C2	121.2 (4)	O3—C11—C12	116.3 (5)
N1—C3—C4	122.6 (3)	C11—C12—H12A	109.5
C2—C3—C4	116.2 (4)	C11—C12—H12B	109.5
C5—C4—C3	121.4 (3)	H12A—C12—H12B	109.5
C5—C4—H4	119.3	C11—C12—H12C	109.5
C3—C4—H4	119.3	H12A—C12—H12C	109.5
C4—C5—C6	121.7 (3)	H12B—C12—H12C	109.5
C4—C5—H5	119.2	C13—O5—H5A	109.5
C6—C5—H5	119.2	O4—C13—O5	122.9 (4)
C5—C6—C1	117.0 (3)	O4—C13—C14	125.9 (4)

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C5—C6—C7	120.8 (3)	O5—C13—C14	111.2 (4)
C1—C6—C7	122.2 (3)	C13—C14—H14A	109.5
N2—C7—C6	121.7 (3)	C13—C14—H14B	109.5
N2—C7—H7	119.1	H14A—C14—H14B	109.5
C6—C7—H7	119.1	C13—C14—H14C	109.5
O1—C8—N4	122.7 (3)	H14A—C14—H14C	109.5
O1—C8—N3	118.5 (3)	H14B—C14—H14C	109.5
C7—N2—N3—C8	-178.5 (4)	C3—C4—C5—C6	0.1 (7)
C6—C1—C2—C3	-0.6 (7)	C4—C5—C6—C1	0.4 (6)
C9—N1—C3—C2	0.7 (7)	C4—C5—C6—C7	-179.1 (4)
C10—N1—C3—C2	-171.2 (4)	C2—C1—C6—C5	-0.1 (6)
C9—N1—C3—C4	-178.3 (4)	C2—C1—C6—C7	179.4 (4)
C10—N1—C3—C4	9.8 (7)	N3—N2—C7—C6	178.7 (4)
C1—C2—C3—N1	-178.0 (4)	C5—C6—C7—N2	177.6 (4)
C1—C2—C3—C4	1.0 (7)	C1—C6—C7—N2	-1.9 (6)
N1—C3—C4—C5	178.3 (4)	N2—N3—C8—O1	175.8 (4)
C2—C3—C4—C5	-0.8 (6)	N2—N3—C8—N4	-5.9 (6)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O3—H3 \cdots O2 ⁱ	0.85	1.82	2.658 (6)	170
O5—H5A \cdots O1 ⁱⁱ	0.82	1.78	2.579 (4)	166
N4—H4A \cdots O4 ⁱⁱⁱ	0.87 (4)	2.48 (4)	3.164 (6)	135 (3)
N3—H3A \cdots O1 ^{iv}	0.897 (10)	2.017 (11)	2.913 (5)	178 (4)
N4—H4B \cdots O4 ^v	0.91 (6)	2.16 (6)	3.051 (6)	166 (5)

Symmetry codes: (i) $-x+1, -y+1, -z+1$; (ii) $x+1, y, z$; (iii) $-x+1, -y+2, -z+1$; (iv) $-x, -y+1, -z+1$; (v) $x-1, y, z$.

Fig. 1

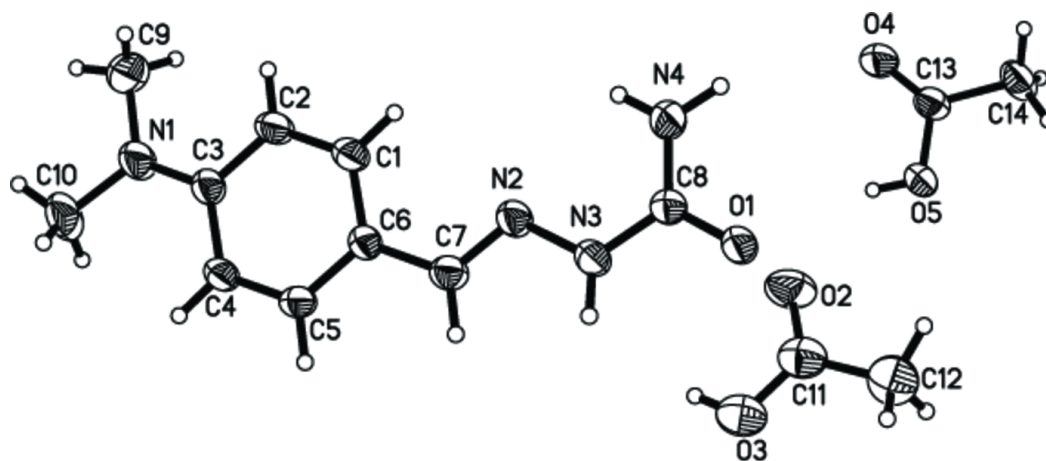


Fig. 2

