

2-Methyl-7-nitro-2,3-dihydro-1-benzofuran

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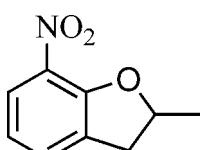
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; R factor = 0.066; wR factor = 0.183; data-to-parameter ratio = 13.1.

The dihydrofuran ring of the title compound, $\text{C}_9\text{H}_9\text{NO}_3$, adopts an envelope conformation. The nitro group is twisted slightly away from the attached benzene ring [dihedral angle = 21.9 (1)°].

Related literature

For bond-length data, see: Allen *et al.* (1987). For details of the synthesis, see: Majumdar *et al.* (2008).



Experimental

Crystal data

$\text{C}_9\text{H}_9\text{NO}_3$	$V = 1722.6$ (6) Å ³
$M_r = 179.17$	$Z = 8$
Orthorhombic, $Pbca$	Mo $K\alpha$ radiation
$a = 8.4250$ (17) Å	$\mu = 0.10$ mm ⁻¹
$b = 7.2260$ (14) Å	$T = 298$ (2) K
$c = 28.295$ (6) Å	$0.30 \times 0.20 \times 0.20$ mm

Data collection

Enraf–Nonius CAD-4 diffractometer	1551 independent reflections
Absorption correction: ψ scan (North <i>et al.</i> , 1968)	829 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.969$, $T_{\max} = 0.979$	$R_{\text{int}} = 0.048$
2977 measured reflections	3 standard reflections
	every 200 reflections
	intensity decay: none

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.066$	118 parameters
$wR(F^2) = 0.182$	H-atom parameters constrained
$S = 1.01$	$\Delta\rho_{\max} = 0.32$ e Å ⁻³
1551 reflections	$\Delta\rho_{\min} = -0.26$ e Å ⁻³

Data collection: *CAD-4 Software* (Enraf–Nonius, 1985); cell refinement: *CAD-4 Software*; 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*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI2613).

References

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supporting information

Acta Cryst. (2008). E64, o1281 [doi:10.1107/S1600536808017728]

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S1. Comment

The title compound, 2-methyl-7-nitro-2,3-dihydrobenzofuran, is an important intermediate for the synthesis of 2-methyl-2,3-dihydrobenzofuran-7-amine. we report here the crystal structure of the title compound.

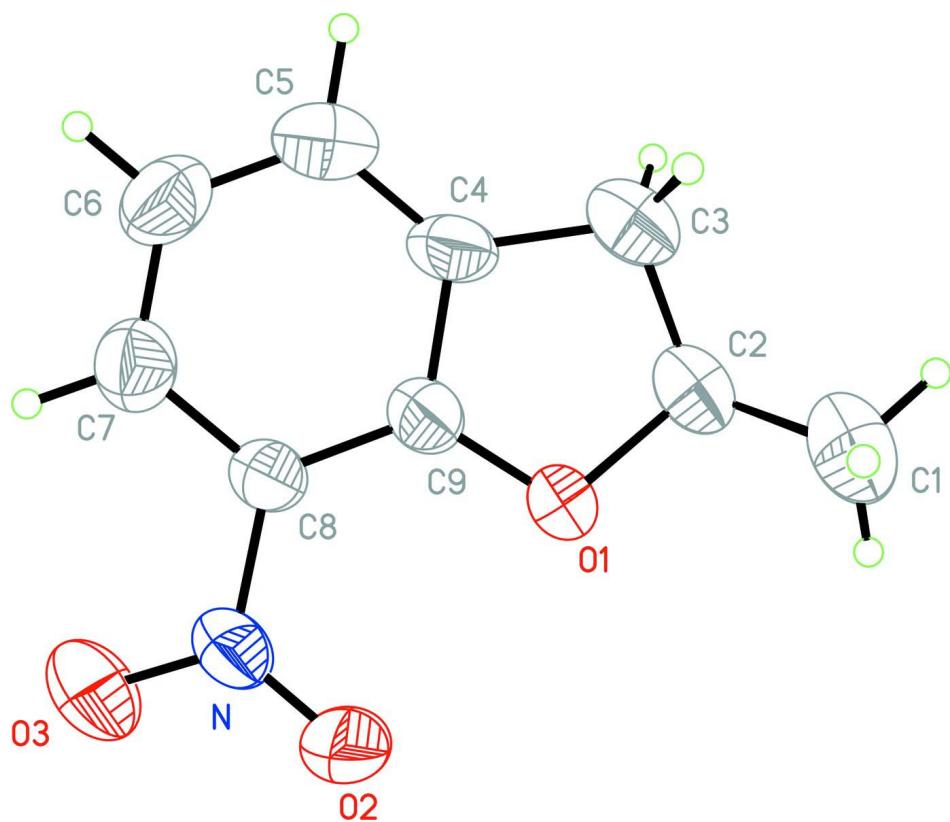
The molecular structure of the compound is shown in Fig. 1. Bond lengths and angles are within normal ranges (Allen *et al.*, 1987), except the C1—C2 bond length of 1.420 (6) Å. The dihydrofuran ring is in an envelope conformation with C2 as flap atom. The nitro group is slightly twisted away from the attached benzene ring [O2—N—C8—C9 = 5.3 (5)° and O3—N—C8—C7 = 5.9 (5)°]. No hydrogen bonding interactions are observed in the crystal structure (Fig.2).

S2. Experimental

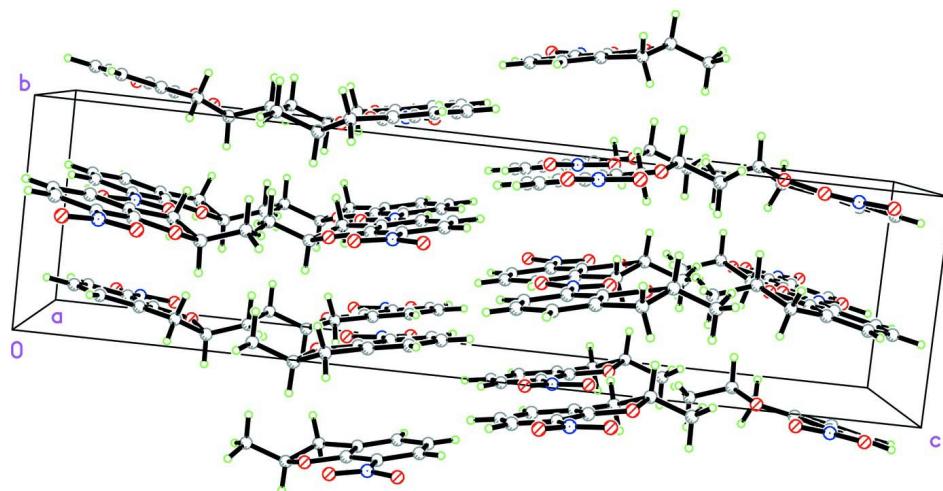
The title compound was synthesized according to the literature method (Majumdar *et al.*, 2008). Single crystals were obtained by slow evaporation of a methanol (25 ml) solution of the compound (0.30 g, 1.6 mmol) at room temperature for about 4 d.

S3. Refinement

H atoms were positioned geometrically [C-H = 0.93-0.98 Å] and constrained to ride on their parent atoms, with $U_{\text{iso}}(\text{H}) = xU_{\text{eq}}(\text{C})$, where $x = 1.2$ for aromatic and methylene H and 1.5 for methyl H atoms.

**Figure 1**

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

**Figure 2**

A view of the molecular packing in the title compound.

2-Methyl-7-nitro-2,3-dihydro-1-benzofuran*Crystal data*

$C_9H_9NO_3$
 $M_r = 179.17$
Orthorhombic, $Pbca$
Hall symbol: -P 2ac 2ab
 $a = 8.4250$ (17) Å
 $b = 7.2260$ (14) Å
 $c = 28.295$ (6) Å
 $V = 1722.6$ (6) Å³
 $Z = 8$

$F(000) = 752$
 $D_x = 1.382$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 25 reflections
 $\theta = 10\text{--}13^\circ$
 $\mu = 0.11$ mm⁻¹
 $T = 298$ K
Block, colourless
0.30 × 0.20 × 0.20 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)

$T_{\min} = 0.969$, $T_{\max} = 0.979$

2977 measured reflections

1551 independent reflections
829 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$
 $\theta_{\max} = 25.2^\circ$, $\theta_{\min} = 1.4^\circ$
 $h = 0 \rightarrow 10$
 $k = -8 \rightarrow 8$
 $l = 0 \rightarrow 33$
3 standard reflections every 200 reflections
intensity decay: none

Refinement

Refinement on F^2

Least-squares matrix: full

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

$wR(F^2) = 0.182$

$S = 1.01$

1551 reflections

118 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
 $w = 1/[\sigma^2(F_o^2) + (0.06P)^2 + 1.5P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.32$ e Å⁻³
 $\Delta\rho_{\min} = -0.26$ e Å⁻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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^* / U_{\text{eq}}$
N	0.5181 (3)	0.0623 (5)	0.39917 (10)	0.0581 (8)
O1	0.2624 (3)	0.0149 (4)	0.32787 (7)	0.0601 (8)
C1	0.1077 (6)	0.0403 (8)	0.25775 (16)	0.0970 (16)
H1A	0.1909	-0.0204	0.2405	0.146*

H1B	0.1258	0.1715	0.2576	0.146*
H1C	0.0074	0.0140	0.2431	0.146*
O2	0.5612 (3)	0.0409 (5)	0.35839 (9)	0.0867 (10)
C2	0.1064 (4)	-0.0249 (8)	0.30507 (14)	0.0812 (14)
H2A	0.0947	-0.1598	0.3038	0.097*
O3	0.6098 (3)	0.0689 (6)	0.43222 (10)	0.0992 (12)
C3	-0.0172 (4)	0.0473 (6)	0.33963 (14)	0.0706 (11)
H3A	-0.0614	0.1638	0.3289	0.085*
H3B	-0.1025	-0.0413	0.3438	0.085*
C4	0.0753 (4)	0.0725 (5)	0.38480 (12)	0.0532 (9)
C5	0.0285 (4)	0.1112 (5)	0.42971 (13)	0.0634 (10)
H5A	-0.0788	0.1240	0.4368	0.076*
C6	0.1416 (5)	0.1314 (6)	0.46479 (13)	0.0624 (10)
H6A	0.1093	0.1557	0.4956	0.075*
C7	0.3000 (4)	0.1164 (5)	0.45498 (12)	0.0560 (9)
H7A	0.3747	0.1319	0.4789	0.067*
C8	0.3491 (4)	0.0774 (5)	0.40863 (11)	0.0460 (8)
C9	0.2360 (4)	0.0537 (4)	0.37377 (11)	0.0459 (8)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N	0.0376 (15)	0.085 (2)	0.0513 (17)	-0.0068 (16)	-0.0090 (16)	0.0044 (16)
O1	0.0358 (11)	0.104 (2)	0.0411 (12)	0.0006 (13)	-0.0058 (12)	-0.0061 (12)
C1	0.066 (3)	0.151 (5)	0.074 (3)	-0.004 (3)	-0.022 (3)	0.017 (3)
O2	0.0389 (13)	0.168 (3)	0.0530 (16)	0.0016 (17)	0.0045 (13)	-0.0016 (17)
C2	0.047 (2)	0.134 (4)	0.063 (2)	-0.004 (2)	-0.014 (2)	-0.002 (3)
O3	0.0446 (15)	0.176 (3)	0.0773 (19)	-0.0046 (19)	-0.0248 (16)	-0.009 (2)
C3	0.0399 (19)	0.098 (3)	0.074 (3)	-0.003 (2)	-0.009 (2)	-0.002 (2)
C4	0.0313 (16)	0.070 (2)	0.058 (2)	0.0012 (16)	0.0062 (16)	0.0053 (18)
C5	0.044 (2)	0.069 (3)	0.077 (3)	0.0068 (18)	0.009 (2)	0.004 (2)
C6	0.069 (2)	0.073 (2)	0.0458 (19)	0.006 (2)	0.011 (2)	-0.0021 (18)
C7	0.060 (2)	0.061 (2)	0.047 (2)	-0.0053 (19)	-0.0032 (18)	0.0004 (17)
C8	0.0352 (17)	0.060 (2)	0.0426 (18)	-0.0023 (15)	-0.0007 (15)	0.0044 (16)
C9	0.0375 (16)	0.0554 (19)	0.0447 (17)	-0.0043 (15)	-0.0028 (16)	0.0030 (16)

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

N—O3	1.214 (4)	C3—H3A	0.97
N—O2	1.219 (3)	C3—H3B	0.97
N—C8	1.453 (4)	C4—C5	1.360 (5)
O1—C9	1.347 (4)	C4—C9	1.396 (4)
O1—C2	1.492 (4)	C5—C6	1.384 (5)
C1—C2	1.420 (6)	C5—H5A	0.93
C1—H1A	0.96	C6—C7	1.367 (5)
C1—H1B	0.96	C6—H6A	0.93
C1—H1C	0.96	C7—C8	1.404 (4)
C2—C3	1.520 (6)	C7—H7A	0.93

C2—H2A	0.98	C8—C9	1.382 (4)
C3—C4	1.508 (5)		
O3—N—O2	123.0 (3)	C2—C3—H3B	111.1
O3—N—C8	118.6 (3)	H3A—C3—H3B	109.0
O2—N—C8	118.4 (3)	C5—C4—C9	120.7 (3)
C9—O1—C2	108.2 (3)	C5—C4—C3	131.8 (3)
C2—C1—H1A	109.5	C9—C4—C3	107.5 (3)
C2—C1—H1B	109.5	C4—C5—C6	119.5 (3)
H1A—C1—H1B	109.5	C4—C5—H5A	120.3
C2—C1—H1C	109.5	C6—C5—H5A	120.3
H1A—C1—H1C	109.5	C7—C6—C5	121.2 (3)
H1B—C1—H1C	109.5	C7—C6—H6A	119.4
C1—C2—O1	109.7 (4)	C5—C6—H6A	119.4
C1—C2—C3	119.9 (4)	C6—C7—C8	119.6 (3)
O1—C2—C3	105.0 (3)	C6—C7—H7A	120.2
C1—C2—H2A	107.2	C8—C7—H7A	120.2
O1—C2—H2A	107.2	C9—C8—C7	119.2 (3)
C3—C2—H2A	107.2	C9—C8—N	122.3 (3)
C4—C3—C2	103.5 (3)	C7—C8—N	118.4 (3)
C4—C3—H3A	111.1	O1—C9—C8	126.9 (3)
C2—C3—H3A	111.1	O1—C9—C4	113.3 (3)
C4—C3—H3B	111.1	C8—C9—C4	119.8 (3)
C9—O1—C2—C1	145.4 (4)	O2—N—C8—C9	5.3 (5)
C9—O1—C2—C3	15.4 (4)	O3—N—C8—C7	5.9 (5)
C1—C2—C3—C4	-139.3 (4)	O2—N—C8—C7	-175.1 (3)
O1—C2—C3—C4	-15.5 (5)	C2—O1—C9—C8	172.0 (4)
C2—C3—C4—C5	-170.7 (4)	C2—O1—C9—C4	-8.9 (4)
C2—C3—C4—C9	10.9 (5)	C7—C8—C9—O1	-179.7 (3)
C9—C4—C5—C6	-0.2 (6)	N—C8—C9—O1	0.0 (6)
C3—C4—C5—C6	-178.4 (4)	C7—C8—C9—C4	1.3 (5)
C4—C5—C6—C7	1.2 (6)	N—C8—C9—C4	-179.1 (3)
C5—C6—C7—C8	-0.8 (6)	C5—C4—C9—O1	179.8 (3)
C6—C7—C8—C9	-0.4 (5)	C3—C4—C9—O1	-1.6 (4)
C6—C7—C8—N	180.0 (3)	C5—C4—C9—C8	-1.0 (6)
O3—N—C8—C9	-173.7 (3)	C3—C4—C9—C8	177.6 (3)