

4,4-Dimethyl-1-(3-nitrophenyl)pent-1-en-3-one

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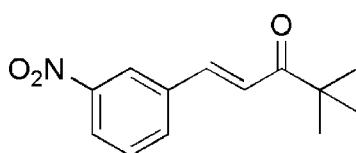
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Key indicators: single-crystal X-ray study; $T = 173\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.041; wR factor = 0.123; data-to-parameter ratio = 12.7.

All the non-hydrogen atoms except one methyl C atom of the title compound, $\text{C}_{13}\text{H}_{15}\text{NO}_3$, lie on a crystallographic mirror plane perpendicular to the b axis. The crystal packing is stabilized by two weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

The title compound is an important intermediate in the pesticides industry (Wang *et al.*, 2006). For related structures, see: Anuradha *et al.* (2008); Butcher *et al.* (2007); Gong *et al.* (2008); Harrison *et al.* (2007); Patil *et al.* (2007); Sarojini *et al.* (2007); Thiruvalluvar *et al.* (2007, 2008); Xia & Hu (2008).



Experimental

Crystal data

$\text{C}_{13}\text{H}_{15}\text{NO}_3$	$V = 1221.72(17)\text{ \AA}^3$
$M_r = 233.26$	$Z = 4$
Orthorhombic, $Pnma$	$\text{Mo K}\alpha$ radiation
$a = 11.3375(9)\text{ \AA}$	$\mu = 0.09\text{ mm}^{-1}$
$b = 7.2163(6)\text{ \AA}$	$T = 173\text{ K}$
$c = 14.9327(12)\text{ \AA}$	$0.48 \times 0.36 \times 0.15\text{ mm}$

Data collection

Bruker SMART 1000 CCD diffractometer	5485 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 2004)	1280 independent reflections
$T_{\min} = 0.958$, $T_{\max} = 0.987$	937 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.030$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$	101 parameters
$wR(F^2) = 0.123$	H-atom parameters constrained
$S = 1.06$	$\Delta\rho_{\text{max}} = 0.26\text{ e \AA}^{-3}$
1280 reflections	$\Delta\rho_{\text{min}} = -0.16\text{ e \AA}^{-3}$

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

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
C6—H6A \cdots O2 ⁱ	0.98	2.44	3.367 (2)	158
C10—H10 \cdots O2 ⁱⁱ	0.95	2.50	3.366 (3)	152

Symmetry codes: (i) $-x + \frac{3}{2}, -y, z + \frac{1}{2}$; (ii) $x - \frac{1}{2}, -y + \frac{1}{2}, -z - \frac{1}{2}$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 2003); data reduction: *SAINT-Plus*; 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: BT2958).

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supporting information

Acta Cryst. (2009). E65, o1346 [doi:10.1107/S1600536809018479]

4,4-Dimethyl-1-(3-nitrophenyl)pent-1-en-3-one

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S1. Comment

The title compound, 1-(3-nitrophenyl)-4,4-dimethylpentan-3-one, is a very important intermediate in the pesticides industry (Wang *et al.*, 2006). Continuing our work (Xia & Hu, 2008), we report the synthesis and crystal structure of the title compound. Several crystal structures containing phenylprop-2-en-1-one moiety have been recently published (Anuradha *et al.*, 2008; Butcher *et al.*, 2007; Gong *et al.*, 2008; Harrison *et al.*, 2007; Patil *et al.*, 2007; Sarojini *et al.*, 2007; Thiruvalluvar *et al.*, 2007, 2008; Xia & Hu, 2008). In the title compound (Fig. 1), the carbonyl, ethenyl and nitro groups are coplanar with the benzene ring. The bond lengths and bond angles in (I) are in excellent agreement with the corresponding bond lengths and angles reported in the related compounds given above. The crystal packing is illustrated in Fig. 2.

S2. Experimental

3,3-dimethylbutan-2-one(0.0105 mol) was added dropwise into the solution of 3-nitrobenzaldehyde (0.01 mol) and 60 ml ethanol. Then 0.1 g 50% NaOH solution as catalyst was added and the solution was stirred at 333 K for 5 h (monitored by TLC). Part of the solvent was evaporated and the solution was cooled to 277 K and a precipitate formed. It was filtered and dried. Crystals suitable for X-ray structure determination were obtained by slow evaporation of an ethanol solution at room temperature.

S3. Refinement

H atoms were positioned geometrically ($C—H = 0.95$ or $C_{methyl}—H = 0.98 \text{ \AA}$) with $U_{iso}(H) = 1.2U_{eq}(C)$ or $1.5U_{eq}(C_{methyl})$. The methyl group on a general position was allowed to rotate but not to tip.

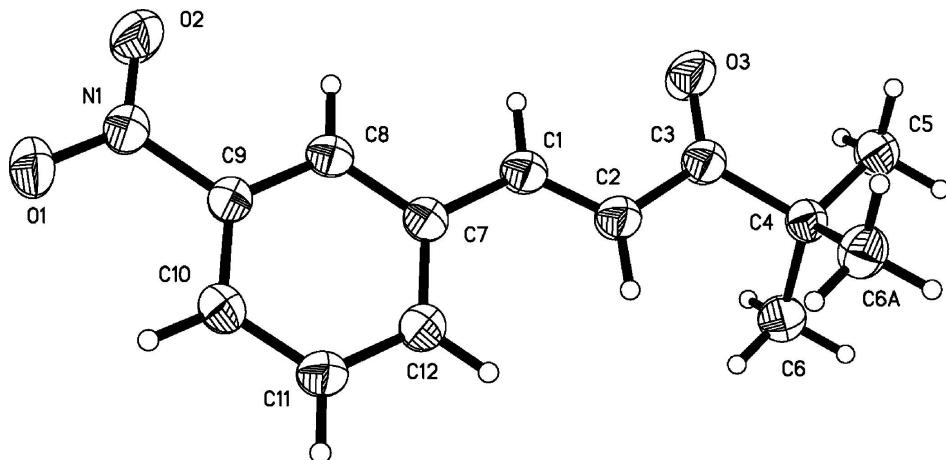
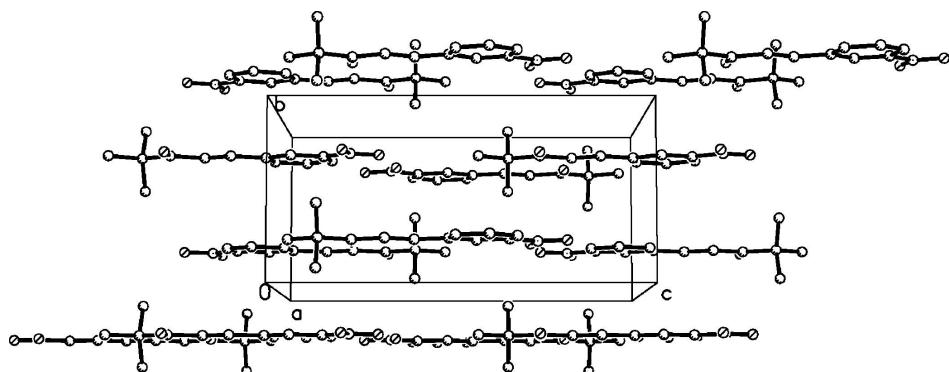


Figure 1

Molecular structure of the title compound, showing 30% probability displacement ellipsoids.

**Figure 2**

Packing diagram of the title compound.

4,4-Dimethyl-1-(3-nitrophenoxy)pent-1-en-3-one

Crystal data

$C_{13}H_{15}NO_3$
 $M_r = 233.26$
 Orthorhombic, $Pnma$
 Hall symbol: -P 2ac 2n
 $a = 11.3375 (9)$ Å
 $b = 7.2163 (6)$ Å
 $c = 14.9327 (12)$ Å
 $V = 1221.72 (17)$ Å³
 $Z = 4$
 $F(000) = 496$

$D_x = 1.268$ Mg m⁻³
 Melting point: 363 K
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 1920 reflections
 $\theta = 2.3\text{--}27.7^\circ$
 $\mu = 0.09$ mm⁻¹
 $T = 173$ K
 Block, colourless
 $0.48 \times 0.36 \times 0.15$ mm

Data collection

Bruker SMART 1000 CCD
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 ω scans
 Absorption correction: multi-scan
 (*SADABS*; Sheldrick, 2004)
 $T_{\min} = 0.958$, $T_{\max} = 0.987$

5485 measured reflections
 1280 independent reflections
 937 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$
 $\theta_{\max} = 26.0^\circ$, $\theta_{\min} = 2.3^\circ$
 $h = -13 \rightarrow 12$
 $k = -3 \rightarrow 8$
 $l = -18 \rightarrow 17$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.123$
 $S = 1.06$
 1280 reflections
 101 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
 $w = 1/[\sigma^2(F_o^2) + (0.0591P)^2 + 0.3634P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.26$ e Å⁻³
 $\Delta\rho_{\min} = -0.16$ 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 (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	U_{iso}^* / U_{eq}	Occ. (<1)
C1	0.64133 (19)	0.2500	0.12364 (15)	0.0363 (6)	
H1	0.7249	0.2500	0.1281	0.044*	
C2	0.5825 (2)	0.2500	0.19956 (15)	0.0371 (6)	
H2	0.4987	0.2500	0.1983	0.045*	
C3	0.6446 (2)	0.2500	0.28774 (15)	0.0367 (6)	
C4	0.56797 (18)	0.2500	0.37208 (15)	0.0317 (5)	
C5	0.6474 (2)	0.2500	0.45437 (16)	0.0392 (6)	
H5A	0.5987	0.2500	0.5086	0.059*	
H5B	0.6973	0.3609	0.4537	0.059*	0.50
H5C	0.6973	0.1391	0.4537	0.059*	0.50
C6	0.48919 (15)	0.0777 (3)	0.37212 (12)	0.0419 (5)	
H6A	0.5385	-0.0338	0.3701	0.063*	
H6B	0.4374	0.0803	0.3196	0.063*	
H6C	0.4412	0.0761	0.4267	0.063*	
C7	0.59173 (19)	0.2500	0.03248 (15)	0.0310 (5)	
C8	0.66910 (19)	0.2500	-0.03967 (15)	0.0313 (5)	
H8	0.7519	0.2500	-0.0299	0.038*	
C9	0.62418 (18)	0.2500	-0.12592 (14)	0.0296 (5)	
C10	0.5051 (2)	0.2500	-0.14355 (15)	0.0348 (6)	
H10	0.4764	0.2500	-0.2033	0.042*	
C11	0.42813 (19)	0.2500	-0.07129 (16)	0.0381 (6)	
H11	0.3454	0.2500	-0.0814	0.046*	
C12	0.47098 (19)	0.2500	0.01538 (15)	0.0363 (6)	
H12	0.4171	0.2500	0.0641	0.044*	
N1	0.70773 (17)	0.2500	-0.20150 (12)	0.0359 (5)	
O1	0.66864 (16)	0.2500	-0.27777 (11)	0.0501 (5)	
O2	0.81309 (15)	0.2500	-0.18457 (12)	0.0570 (6)	
O3	0.75145 (14)	0.2500	0.29171 (11)	0.0559 (6)	

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0277 (11)	0.0469 (15)	0.0344 (13)	0.000	-0.0032 (9)	0.000
C2	0.0283 (11)	0.0495 (15)	0.0337 (13)	0.000	-0.0012 (9)	0.000
C3	0.0319 (12)	0.0453 (15)	0.0328 (13)	0.000	-0.0006 (9)	0.000
C4	0.0315 (11)	0.0347 (13)	0.0290 (12)	0.000	-0.0002 (9)	0.000

C5	0.0373 (12)	0.0471 (15)	0.0333 (13)	0.000	-0.0019 (10)	0.000
C6	0.0415 (9)	0.0424 (10)	0.0418 (10)	-0.0048 (8)	0.0023 (7)	0.0001 (8)
C7	0.0304 (11)	0.0317 (13)	0.0310 (12)	0.000	-0.0015 (9)	0.000
C8	0.0265 (10)	0.0332 (13)	0.0340 (13)	0.000	-0.0030 (9)	0.000
C9	0.0312 (11)	0.0281 (12)	0.0297 (12)	0.000	0.0014 (9)	0.000
C10	0.0352 (12)	0.0387 (14)	0.0306 (12)	0.000	-0.0043 (9)	0.000
C11	0.0279 (11)	0.0494 (15)	0.0372 (13)	0.000	-0.0014 (9)	0.000
C12	0.0321 (12)	0.0447 (14)	0.0322 (13)	0.000	0.0006 (9)	0.000
N1	0.0355 (11)	0.0405 (12)	0.0316 (12)	0.000	0.0020 (8)	0.000
O1	0.0501 (10)	0.0725 (14)	0.0278 (10)	0.000	-0.0010 (8)	0.000
O2	0.0322 (9)	0.0955 (16)	0.0434 (11)	0.000	0.0058 (8)	0.000
O3	0.0300 (10)	0.1027 (17)	0.0350 (10)	0.000	-0.0011 (7)	0.000

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

C1—C2	1.315 (3)	C6—H6C	0.9800
C1—C7	1.473 (3)	C7—C8	1.389 (3)
C1—H1	0.9500	C7—C12	1.393 (3)
C2—C3	1.493 (3)	C8—C9	1.385 (3)
C2—H2	0.9500	C8—H8	0.9500
C3—O3	1.213 (3)	C9—C10	1.376 (3)
C3—C4	1.530 (3)	C9—N1	1.474 (3)
C4—C5	1.523 (3)	C10—C11	1.388 (3)
C4—C6	1.531 (2)	C10—H10	0.9500
C4—C6 ⁱ	1.531 (2)	C11—C12	1.382 (3)
C5—H5A	0.9800	C11—H11	0.9500
C5—H5B	0.9800	C12—H12	0.9500
C5—H5C	0.9800	N1—O2	1.221 (2)
C6—H6A	0.9800	N1—O1	1.222 (2)
C6—H6B	0.9800		
C2—C1—C7	127.1 (2)	C4—C6—H6C	109.5
C2—C1—H1	116.5	H6A—C6—H6C	109.5
C7—C1—H1	116.5	H6B—C6—H6C	109.5
C1—C2—C3	121.4 (2)	C8—C7—C12	118.6 (2)
C1—C2—H2	119.3	C8—C7—C1	118.40 (19)
C3—C2—H2	119.3	C12—C7—C1	123.0 (2)
O3—C3—C2	120.9 (2)	C9—C8—C7	119.27 (19)
O3—C3—C4	121.8 (2)	C9—C8—H8	120.4
C2—C3—C4	117.30 (19)	C7—C8—H8	120.4
C5—C4—C3	109.19 (18)	C10—C9—C8	122.6 (2)
C5—C4—C6	110.15 (12)	C10—C9—N1	118.97 (19)
C3—C4—C6	109.36 (12)	C8—C9—N1	118.42 (18)
C5—C4—C6 ⁱ	110.15 (12)	C9—C10—C11	117.9 (2)
C3—C4—C6 ⁱ	109.36 (12)	C9—C10—H10	121.0
C6—C4—C6 ⁱ	108.63 (18)	C11—C10—H10	121.0
C4—C5—H5A	109.5	C12—C11—C10	120.5 (2)
C4—C5—H5B	109.5	C12—C11—H11	119.8

H5A—C5—H5B	109.5	C10—C11—H11	119.8
C4—C5—H5C	109.5	C11—C12—C7	121.1 (2)
H5A—C5—H5C	109.5	C11—C12—H12	119.4
H5B—C5—H5C	109.5	C7—C12—H12	119.4
C4—C6—H6A	109.5	O2—N1—O1	123.22 (19)
C4—C6—H6B	109.5	O2—N1—C9	118.06 (18)
H6A—C6—H6B	109.5	O1—N1—C9	118.73 (18)
C7—C1—C2—C3	180.0	C7—C8—C9—C10	0.0
C1—C2—C3—O3	0.0	C7—C8—C9—N1	180.0
C1—C2—C3—C4	180.0	C8—C9—C10—C11	0.0
O3—C3—C4—C5	0.0	N1—C9—C10—C11	180.0
C2—C3—C4—C5	180.0	C9—C10—C11—C12	0.0
O3—C3—C4—C6	−120.58 (12)	C10—C11—C12—C7	0.0
C2—C3—C4—C6	59.42 (12)	C8—C7—C12—C11	0.0
O3—C3—C4—C6 ⁱ	120.58 (12)	C1—C7—C12—C11	180.0
C2—C3—C4—C6 ⁱ	−59.42 (12)	C10—C9—N1—O2	180.0
C2—C1—C7—C8	180.0	C8—C9—N1—O2	0.0
C2—C1—C7—C12	0.0	C10—C9—N1—O1	0.0
C12—C7—C8—C9	0.0	C8—C9—N1—O1	180.0
C1—C7—C8—C9	180.0		

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

Hydrogen-bond geometry (\AA , °)

$D\text{—H}\cdots A$	$D\text{—H}$	$\text{H}\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
C6—H6A ⁱⁱ —O2 ⁱⁱ	0.98	2.44	3.367 (2)	158
C10—H10 ⁱⁱⁱ —O2 ⁱⁱⁱ	0.95	2.50	3.366 (3)	152

Symmetry codes: (ii) $-x+3/2, -y, z+1/2$; (iii) $x-1/2, -y+1/2, -z-1/2$.