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2-Iodo-3-(4-methoxyanilino)-5,5-dimethylcyclohex-2-en-1-one

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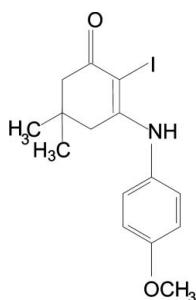
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.027; wR factor = 0.074; data-to-parameter ratio = 22.0.

The cyclohexene ring in the title compound, $\text{C}_{15}\text{H}_{18}\text{INO}_2$, adopts a sofa conformation. The dihedral angle between the cyclohexene (through all ring atoms) and benzene rings is $63.3(1)^\circ$. The molecular conformation features an $\text{N}-\text{H}\cdots\text{I}$ short contact and the crystal packing features $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds.

Related literature

For the biological activity of cyclohex-2-enone derivatives, see: Correia *et al.* (2001); Rebacz *et al.* (2007); Stadler *et al.* (1994). For the use of cyclohex-2-enone in organic synthesis, see: Cokcer *et al.* (1995); Pandey *et al.* (2004). For pukering parameters, see: Cremer & Pople, (1975). For related structures, see: Mohan *et al.* (2008); North *et al.* (2011).



Experimental

Crystal data

 $\text{C}_{15}\text{H}_{18}\text{INO}_2$
 $M_r = 371.20$ Orthorhombic, $Pbca$
 $a = 15.922(5)$ Å $b = 10.107(5)$ Å
 $c = 19.034(5)$ Å
 $V = 3063(2)$ Å³
 $Z = 8$ Mo $K\alpha$ radiation
 $\mu = 2.09$ mm⁻¹
 $T = 298$ K
 $0.20 \times 0.20 \times 0.20$ mm

Data collection

Bruker SMART APEXII area-
detector diffractometer
15382 measured reflections3785 independent reflections
2793 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.027$
 $wR(F^2) = 0.074$
 $S = 0.93$
3785 reflections172 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.46$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.53$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C}14-\text{H}14\cdots\text{O}1^i$	0.93	2.39	3.313 (3)	174
$\text{N}1-\text{H}1\cdots\text{I}1$	0.86	2.71	3.227 (2)	120

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

Data collection: *APEX2* (Bruker, 2008); cell refinement: *SAINT* (Bruker, 2008); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97*, *PLATON* and *pubCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KP2383).

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supporting information

Acta Cryst. (2012). E68, o506 [doi:10.1107/S1600536812002255]

2-Iodo-3-(4-methoxyanilino)-5,5-dimethylcyclohex-2-en-1-one

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S1. Comment

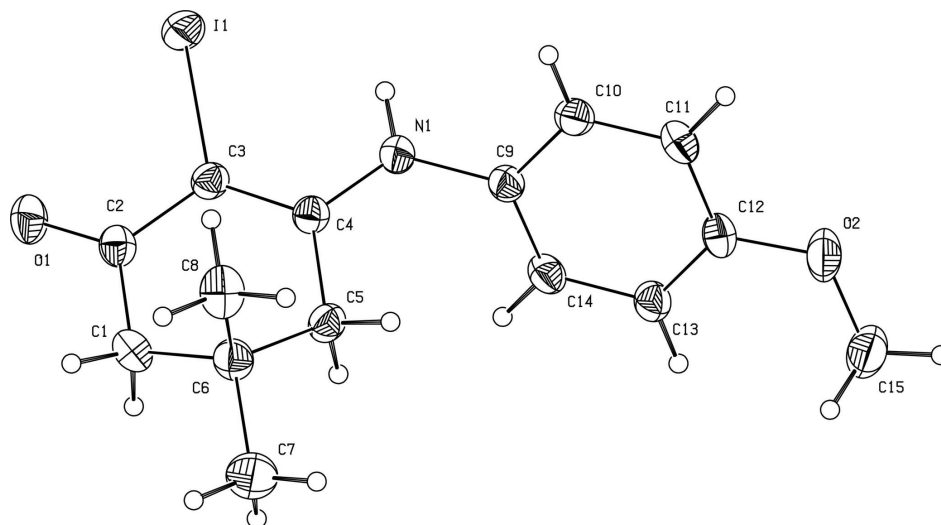
Cyclohex-2-enone derivative exhibits antibacterial (Stadler *et al.*, 1994) and anticancer (Correia *et al.*, 2001; Rebacz *et al.*, 2007) activities. Cyclohex-2-enone plays an important role in organic synthesis (Cokcer *et al.*, 1995; Pandey *et al.*, 2004). Against this background, the title compound was chosen for X-ray structure analysis (Fig. 1). The cyclohexene ring adopts a sofa conformation with the puckering parameters (Cremer & Pople, 1975) being $q_2=0.409$ (3) Å, $q_3=-0.247$ (3) Å and $Q_T=0.478$ (3) Å. The molecular structure is stabilised by N—H···I intramolecular interactions and the crystal packing is stabilised by C—H···O hydrogen bonds (Fig. 2 and Table 1).

S2. Experimental

1,3-cyclohexanedione (2 mmol), FeCl₃.6H₂O (5 mol), and 50 mg of sodium sulfate were successively added in a dry Schlenk tube under argon. The solids were then dissolved in 3 mL of dichloromethane and stirred for 5 m. Aniline (2 mmol) was slowly added and the dark brown coloured mixture was allowed to stir overnight. After completion, solvents were removed under vacuum and the crude oil was filtered on a plug of neutral alumina (eluent: dichloromethane/methanol, 90/10). Solvents were then removed and enaminone product was obtained as a bright yellow solid. Then iodine (3 mmol) dissolved in CCl₄/pyridine (1;1, 10 mL) was added dropwise under an atmosphere of argon to a solution of enaminone (1.5 mmol) in CCl₄/pyridine (1;1, 10 mL) at 273 K. The mixture was stirred for 2 h during that time the temperature was allowed to raise to room temperature. The mixture was diluted with ethyl acetate (50 mL) and washed successively with 1 NHCl (4 × 10 mL), sat. NaHCO₃ (20 mL), 20% aqueous Na₂S₂O₃ (20 mL) and dried (Na₂SO₄). Filtered and concentrated under reduced pressure, the residue was further purified by column chromatography to afford pure 2-iodo-5,5-dimethyl-3-(Phenylamino) cyclohex-2-enone.

S3. Refinement

Hydrogen atoms were positioned geometrically and allowed to ride on their parent atoms, with C—H = 0.93 - 0.97 Å and $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms and 1.2 $U_{eq}(C)$ for other H atoms.

**Figure 1**

The molecular structure of the title compound showing the atom-numbering scheme and displacement ellipsoids drawn at the 30% probability level.

Figure 2

The crystal packing of the title compound. Hydrogen bonds are shown by dashed lines.

2-Iodo-3-(4-methoxyanilino)-5,5-dimethylcyclohex-2-en-1-one

Crystal data

$C_{15}H_{18}INO_2$

$M_r = 371.20$

Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

$a = 15.922 (5) \text{ \AA}$

$b = 10.107 (5) \text{ \AA}$

$c = 19.034 (5) \text{ \AA}$

$V = 3063 (2) \text{ \AA}^3$

$Z = 8$

$F(000) = 1472$

$D_x = 1.610 \text{ Mg m}^{-3}$

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

Cell parameters from 3785 reflections

$\theta = 2.1\text{--}28.3^\circ$

$\mu = 2.09 \text{ mm}^{-1}$

$T = 298 \text{ K}$

Block, colourless

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

Data collection

Bruker SMART APEXII area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω and φ scans

15382 measured reflections

3785 independent reflections

2793 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.025$

$\theta_{\text{max}} = 28.3^\circ$, $\theta_{\text{min}} = 2.1^\circ$

$h = -20 \rightarrow 21$

$k = -13 \rightarrow 13$

$l = -18 \rightarrow 25$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.027$

$wR(F^2) = 0.074$

$S = 0.93$

3785 reflections

172 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

$$w = 1/[\sigma^2(F_o^2) + (0.0379P)^2 + 1.6353P]$$

where $P = (F_o^2 + 2F_c^2)/3$

$$(\Delta/\sigma)_{\max} = 0.001$$

$$\Delta\rho_{\max} = 0.46 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.53 \text{ e } \text{\AA}^{-3}$$

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
I1	0.423501 (11)	0.337299 (16)	0.478996 (11)	0.05300 (8)
O1	0.52198 (11)	0.14887 (18)	0.37743 (10)	0.0540 (5)
O2	0.07995 (12)	-0.1517 (2)	0.71999 (11)	0.0631 (6)
N1	0.30564 (14)	0.1095 (2)	0.54533 (12)	0.0460 (5)
H1	0.3092	0.1935	0.5520	0.055*
C1	0.45213 (16)	-0.0582 (3)	0.38180 (14)	0.0477 (6)
H1A	0.4938	-0.1126	0.4052	0.057*
H1B	0.4621	-0.0650	0.3317	0.057*
C2	0.46504 (15)	0.0831 (2)	0.40367 (13)	0.0401 (5)
C3	0.40963 (15)	0.1356 (2)	0.45650 (14)	0.0390 (5)
C4	0.35334 (15)	0.0597 (2)	0.49300 (12)	0.0378 (5)
C5	0.34440 (16)	-0.0847 (2)	0.47456 (12)	0.0417 (5)
H5A	0.2873	-0.1125	0.4844	0.050*
H5B	0.3816	-0.1360	0.5044	0.050*
C6	0.36464 (17)	-0.1148 (3)	0.39783 (13)	0.0456 (6)
C7	0.3655 (2)	-0.2658 (3)	0.38817 (17)	0.0698 (9)
H7A	0.3780	-0.2866	0.3401	0.105*
H7B	0.3115	-0.3012	0.4003	0.105*
H7C	0.4076	-0.3039	0.4181	0.105*
C8	0.29981 (18)	-0.0539 (3)	0.34872 (15)	0.0585 (7)
H8A	0.3142	-0.0742	0.3009	0.088*
H8B	0.2990	0.0403	0.3550	0.088*
H8C	0.2453	-0.0895	0.3592	0.088*
C9	0.24994 (15)	0.0393 (2)	0.59096 (12)	0.0381 (5)
C10	0.16832 (16)	0.0851 (2)	0.59962 (13)	0.0424 (5)
H10	0.1502	0.1594	0.5751	0.051*
C11	0.11454 (15)	0.0208 (3)	0.64420 (13)	0.0455 (6)
H11	0.0605	0.0536	0.6508	0.055*
C12	0.13951 (16)	-0.0923 (3)	0.67950 (12)	0.0437 (6)
C13	0.22144 (16)	-0.1371 (3)	0.67234 (13)	0.0437 (6)

H13	0.2393	-0.2119	0.6965	0.052*
C14	0.27628 (15)	-0.0696 (2)	0.62894 (13)	0.0432 (5)
H14	0.3317	-0.0978	0.6252	0.052*
C15	0.1004 (2)	-0.2714 (3)	0.75369 (18)	0.0698 (9)
H15A	0.0528	-0.3020	0.7801	0.105*
H15B	0.1469	-0.2573	0.7849	0.105*
H15C	0.1154	-0.3364	0.7191	0.105*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
I1	0.05114 (12)	0.03804 (11)	0.06982 (15)	-0.00787 (7)	0.00678 (9)	-0.00135 (8)
O1	0.0359 (9)	0.0657 (12)	0.0603 (12)	-0.0075 (8)	0.0092 (9)	0.0037 (9)
O2	0.0516 (12)	0.0801 (16)	0.0577 (12)	-0.0013 (10)	0.0190 (9)	0.0170 (10)
N1	0.0504 (12)	0.0374 (11)	0.0503 (12)	-0.0041 (9)	0.0158 (10)	-0.0014 (9)
C1	0.0425 (13)	0.0544 (15)	0.0463 (14)	0.0056 (12)	0.0054 (12)	-0.0065 (12)
C2	0.0310 (11)	0.0487 (14)	0.0406 (13)	-0.0014 (10)	-0.0019 (10)	0.0016 (11)
C3	0.0388 (12)	0.0349 (12)	0.0432 (13)	-0.0038 (9)	-0.0004 (10)	-0.0004 (10)
C4	0.0368 (12)	0.0384 (12)	0.0381 (12)	-0.0022 (10)	-0.0006 (10)	-0.0005 (10)
C5	0.0459 (14)	0.0373 (12)	0.0419 (13)	-0.0069 (10)	0.0024 (11)	0.0000 (10)
C6	0.0489 (14)	0.0450 (14)	0.0428 (14)	-0.0053 (11)	0.0023 (12)	-0.0077 (11)
C7	0.088 (2)	0.0530 (18)	0.0683 (19)	-0.0100 (17)	0.0057 (18)	-0.0191 (15)
C8	0.0502 (15)	0.075 (2)	0.0503 (16)	-0.0110 (15)	-0.0060 (13)	-0.0034 (14)
C9	0.0406 (12)	0.0369 (11)	0.0370 (12)	-0.0015 (10)	0.0040 (10)	-0.0040 (10)
C10	0.0464 (13)	0.0411 (13)	0.0398 (13)	0.0056 (11)	0.0006 (11)	0.0000 (10)
C11	0.0354 (12)	0.0572 (16)	0.0440 (13)	0.0067 (11)	0.0054 (11)	-0.0043 (12)
C12	0.0440 (13)	0.0552 (15)	0.0318 (12)	-0.0019 (11)	0.0072 (11)	-0.0021 (11)
C13	0.0458 (14)	0.0488 (14)	0.0367 (13)	0.0052 (11)	0.0041 (11)	0.0042 (10)
C14	0.0358 (12)	0.0499 (14)	0.0440 (13)	0.0078 (11)	0.0038 (11)	-0.0008 (11)
C15	0.083 (2)	0.067 (2)	0.0587 (18)	-0.0143 (18)	0.0167 (17)	0.0092 (16)

Geometric parameters (Å, °)

I1—C3	2.095 (3)	C7—H7A	0.9600
O1—C2	1.230 (3)	C7—H7B	0.9600
O2—C12	1.361 (3)	C7—H7C	0.9600
O2—C15	1.407 (4)	C8—H8A	0.9600
N1—C4	1.350 (3)	C8—H8B	0.9600
N1—C9	1.430 (3)	C8—H8C	0.9600
N1—H1	0.8600	C9—C14	1.382 (3)
C1—C2	1.501 (4)	C9—C10	1.389 (3)
C1—C6	1.537 (4)	C10—C11	1.369 (3)
C1—H1A	0.9700	C10—H10	0.9300
C1—H1B	0.9700	C11—C12	1.385 (4)
C2—C3	1.439 (3)	C11—H11	0.9300
C3—C4	1.369 (3)	C12—C13	1.387 (3)
C4—C5	1.508 (3)	C13—C14	1.382 (3)
C5—C6	1.526 (3)	C13—H13	0.9300

C5—H5A	0.9700	C14—H14	0.9300
C5—H5B	0.9700	C15—H15A	0.9600
C6—C8	1.523 (4)	C15—H15B	0.9600
C6—C7	1.537 (4)	C15—H15C	0.9600
C12—O2—C15	118.4 (2)	C6—C7—H7C	109.5
C4—N1—C9	127.7 (2)	H7A—C7—H7C	109.5
C4—N1—H1	116.1	H7B—C7—H7C	109.5
C9—N1—H1	116.1	C6—C8—H8A	109.5
C2—C1—C6	115.0 (2)	C6—C8—H8B	109.5
C2—C1—H1A	108.5	H8A—C8—H8B	109.5
C6—C1—H1A	108.5	C6—C8—H8C	109.5
C2—C1—H1B	108.5	H8A—C8—H8C	109.5
C6—C1—H1B	108.5	H8B—C8—H8C	109.5
H1A—C1—H1B	107.5	C14—C9—C10	119.1 (2)
O1—C2—C3	122.4 (2)	C14—C9—N1	121.7 (2)
O1—C2—C1	120.2 (2)	C10—C9—N1	119.1 (2)
C3—C2—C1	117.4 (2)	C11—C10—C9	120.0 (2)
C4—C3—C2	123.3 (2)	C11—C10—H10	120.0
C4—C3—H1	120.70 (18)	C9—C10—H10	120.0
C2—C3—H1	115.92 (17)	C10—C11—C12	120.9 (2)
N1—C4—C3	122.3 (2)	C10—C11—H11	119.6
N1—C4—C5	118.7 (2)	C12—C11—H11	119.6
C3—C4—C5	119.1 (2)	O2—C12—C11	116.0 (2)
C4—C5—C6	113.3 (2)	O2—C12—C13	124.5 (2)
C4—C5—H5A	108.9	C11—C12—C13	119.4 (2)
C6—C5—H5A	108.9	C14—C13—C12	119.5 (2)
C4—C5—H5B	108.9	C14—C13—H13	120.3
C6—C5—H5B	108.9	C12—C13—H13	120.3
H5A—C5—H5B	107.7	C9—C14—C13	120.9 (2)
C8—C6—C5	111.3 (2)	C9—C14—H14	119.5
C8—C6—C1	110.0 (2)	C13—C14—H14	119.5
C5—C6—C1	107.9 (2)	O2—C15—H15A	109.5
C8—C6—C7	109.5 (2)	O2—C15—H15B	109.5
C5—C6—C7	108.3 (2)	H15A—C15—H15B	109.5
C1—C6—C7	109.7 (2)	O2—C15—H15C	109.5
C6—C7—H7A	109.5	H15A—C15—H15C	109.5
C6—C7—H7B	109.5	H15B—C15—H15C	109.5
H7A—C7—H7B	109.5		
C6—C1—C2—O1	-160.3 (2)	C2—C1—C6—C8	71.8 (3)
C6—C1—C2—C3	20.7 (3)	C2—C1—C6—C5	-49.8 (3)
O1—C2—C3—C4	-171.1 (2)	C2—C1—C6—C7	-167.7 (2)
C1—C2—C3—C4	7.9 (4)	C4—N1—C9—C14	53.2 (4)
O1—C2—C3—H1	6.0 (3)	C4—N1—C9—C10	-129.6 (3)
C1—C2—C3—H1	-174.98 (17)	C14—C9—C10—C11	-1.4 (4)
C9—N1—C4—C3	-174.5 (2)	N1—C9—C10—C11	-178.7 (2)
C9—N1—C4—C5	5.2 (4)	C9—C10—C11—C12	-2.0 (4)

C2—C3—C4—N1	175.8 (2)	C15—O2—C12—C11	176.3 (3)
I1—C3—C4—N1	-1.1 (3)	C15—O2—C12—C13	-4.2 (4)
C2—C3—C4—C5	-3.9 (4)	C10—C11—C12—O2	-177.1 (2)
I1—C3—C4—C5	179.19 (17)	C10—C11—C12—C13	3.3 (4)
N1—C4—C5—C6	151.7 (2)	O2—C12—C13—C14	179.2 (2)
C3—C4—C5—C6	-28.6 (3)	C11—C12—C13—C14	-1.3 (4)
C4—C5—C6—C8	-67.4 (3)	C10—C9—C14—C13	3.4 (4)
C4—C5—C6—C1	53.4 (3)	N1—C9—C14—C13	-179.4 (2)
C4—C5—C6—C7	172.1 (2)	C12—C13—C14—C9	-2.0 (4)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C14—H14...O1 ⁱ	0.93	2.39	3.313 (3)	174
N1—H1...I1	0.86	2.71	3.227 (2)	120

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