

4-(Dimethylamino)phenyl phenyl ketone

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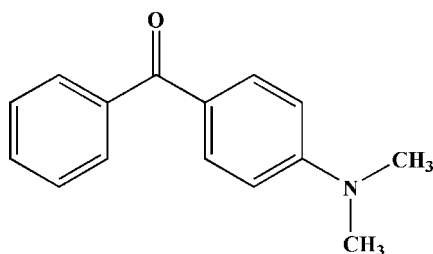
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 Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.001$ Å; R factor = 0.047; wR factor = 0.139; data-to-parameter ratio = 33.1.

In the crystal structure of the title compound, $\text{C}_{15}\text{H}_{15}\text{NO}$, the two benzene rings are twisted from each other by a dihedral angle of $47.97(4)^\circ$. The crystal structure is stabilized by weak intermolecular $\text{C}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\pi$ interactions, and $\pi-\pi$ interactions with a centroid-centroid distance of $3.8493(5)$ Å are observed.

Related literature

For related literature on non-linear optical properties of benzophenone, see: Arivanandhan *et al.* (2006); Szyszyng *et al.* (2004); Vijayan *et al.* (2002) & Wang *et al.*, (2007). For bond-length data see: Allen *et al.* (1987)



Experimental

Crystal data

$\text{C}_{15}\text{H}_{15}\text{NO}$
 $M_r = 225.28$
 Monoclinic, $P2_1/c$
 $a = 13.0575(3)$ Å
 $b = 7.7456(2)$ Å
 $c = 12.4931(3)$ Å
 $\beta = 111.717(1)^\circ$

$V = 1173.85(5)$ Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 100.0(1)$ K
 $0.60 \times 0.43 \times 0.28$ mm

Data collection

Bruker SMART APEXII CCD
 area-detector diffractometer
 Absorption correction: multi-scan
 (SADABS; Bruker, 2005)
 $T_{\min} = 0.939$, $T_{\max} = 0.977$

22302 measured reflections
 5156 independent reflections
 4138 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.138$
 $S = 1.06$
 5156 reflections

156 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.42$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.25$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{C4}-\text{H4}\cdots\text{O1}^i$	0.93	2.46	3.3730 (12)	168
$\text{C10}-\text{H10}\cdots\text{Cg2}^{\text{ii}}$	0.93	2.98	3.6452 (9)	130

Symmetry codes: (i) $-x + 1, y - \frac{1}{2}, -z + \frac{1}{2}$; (ii) $-x, y + \frac{1}{2}, -z + \frac{1}{2}$. Cg2 is the centroid of atoms C8–C13.

Data collection: APEX2 (Bruker, 2005); cell refinement: APEX2; data reduction: SAINT (Bruker, 2005); program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL and PLATON (Spek, 2003).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: AT2585).

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supplementary materials

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4-(Dimethylamino)phenyl phenyl ketone

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Comment

Benzophenone and its derivatives exhibits non-linear optical properties (Wang *et al.*, 2007; Vijayan *et al.*, 2002 & Arivanandhan *et al.*, 2006) and are good candidates for the non-linear optical applications (Szyrszyng *et al.*, 2004). In view of the importance of the benzophenone derivatives, the crystal structure of the title compound (I) has been elucidated.

The asymmetric unit of (I) consists of one molecule of 4-(dimethylamino)benzophenone. Bond lengths and angles in the molecule are found to have normal values (Allen *et al.*, 1987) The dihedral angle formed by the rings (C1–C6) and (C8–C13) is 47.97 (4)° indicating that the rings are twisted from each other. The crystal packing (Fig.2) is consolidated by intermolecular C—H...O hydrogen bonds and C—H... π interactions. π – π interactions with the centroid to centroid distance of 3.8493 (5)Å are observed.

Experimental

4-(Dimethylamino)benzophenone was purchased from Aldrich and dissolved in ethanol. The solution was allowed to evaporate slowly. Colourless crystals were obtained after a month.

Refinement

H atoms were positioned geometrically [C—H = 0.93Å and CH₃=0.96 Å] and refined using a riding model, with $U_{iso}(H)$ = 1.2 or 1.5 $U_{eq}(C)$. The rotating group model was considered for the methyl H atoms.

Figures

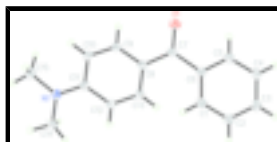


Fig. 1. The molecular structure of the title compound, showing 50% probability displacement ellipsoids and the atom numbering scheme.

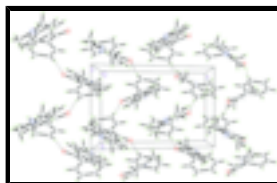


Fig. 2. The crystal packing of the title compound, viewed along the *a* axis. Hydrogen bonds are shown as dashed lines.

4-(Dimethylamino)phenyl phenyl ketone

Crystal data

$C_{15}H_{15}NO$	$F_{000} = 480$
$M_r = 225.28$	$D_x = 1.275 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
Hall symbol: -P 2ybc	$\lambda = 0.71073 \text{ \AA}$
$a = 13.0575 (3) \text{ \AA}$	Cell parameters from 8685 reflections
$b = 7.7456 (2) \text{ \AA}$	$\theta = 3.1\text{--}38.7^\circ$
$c = 12.4931 (3) \text{ \AA}$	$\mu = 0.08 \text{ mm}^{-1}$
$\beta = 111.7170 (10)^\circ$	$T = 100.0 (1) \text{ K}$
$V = 1173.85 (5) \text{ \AA}^3$	Block, colourless
$Z = 4$	$0.60 \times 0.43 \times 0.28 \text{ mm}$

Data collection

Bruker SMART APEXII CCD area-detector diffractometer	5156 independent reflections
Radiation source: fine-focus sealed tube	4138 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\text{int}} = 0.028$
$T = 100.0(1) \text{ K}$	$\theta_{\text{max}} = 35.0^\circ$
φ and ω scans	$\theta_{\text{min}} = 3.1^\circ$
Absorption correction: multi-scan (SADABS; Bruker, 2005)	$h = -21 \rightarrow 19$
$T_{\text{min}} = 0.939$, $T_{\text{max}} = 0.977$	$k = -10 \rightarrow 12$
22302 measured reflections	$l = -20 \rightarrow 20$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.047$	H-atom parameters constrained
$wR(F^2) = 0.138$	$w = 1/[\sigma^2(F_o^2) + (0.0744P)^2 + 0.1876P]$
$S = 1.06$	where $P = (F_o^2 + 2F_c^2)/3$
5156 reflections	$(\Delta/\sigma)_{\text{max}} < 0.001$
156 parameters	$\Delta\rho_{\text{max}} = 0.42 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	$\Delta\rho_{\text{min}} = -0.25 \text{ e \AA}^{-3}$
	Extinction correction: none

Special details

Experimental. The data was collected with the Oxford Cyrosystem Cobra low-temperature attachment.

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.26958 (5)	0.42158 (10)	0.20400 (5)	0.02533 (15)
N1	-0.10640 (5)	0.34279 (10)	0.40740 (6)	0.01934 (14)
C1	0.43352 (6)	0.31796 (11)	0.49440 (7)	0.01861 (15)
H1	0.3933	0.3765	0.5308	0.022*
C2	0.53797 (7)	0.25436 (12)	0.55889 (7)	0.02153 (16)
H2	0.5677	0.2714	0.6382	0.026*
C3	0.59761 (7)	0.16567 (12)	0.50493 (8)	0.02318 (17)
H3	0.6668	0.1215	0.5484	0.028*
C4	0.55457 (7)	0.14235 (12)	0.38612 (8)	0.02373 (17)
H4	0.5948	0.0833	0.3500	0.028*
C5	0.45115 (7)	0.20786 (11)	0.32186 (7)	0.02078 (16)
H5	0.4228	0.1942	0.2423	0.025*
C6	0.38907 (6)	0.29412 (10)	0.37532 (7)	0.01685 (14)
C7	0.27935 (6)	0.36465 (10)	0.29960 (7)	0.01733 (14)
C8	0.18415 (6)	0.36068 (10)	0.33575 (6)	0.01575 (14)
C9	0.08856 (6)	0.44970 (10)	0.26790 (7)	0.01806 (15)
H9	0.0897	0.5141	0.2055	0.022*
C10	-0.00688 (6)	0.44473 (11)	0.29074 (7)	0.01847 (15)
H10	-0.0687	0.5052	0.2436	0.022*
C11	-0.01199 (6)	0.34866 (10)	0.38509 (6)	0.01540 (14)
C12	0.08462 (6)	0.25969 (10)	0.45427 (6)	0.01625 (14)
H12	0.0843	0.1963	0.5174	0.019*
C13	0.17938 (6)	0.26547 (10)	0.42967 (6)	0.01610 (14)
H13	0.2414	0.2051	0.4763	0.019*
C14	-0.11461 (7)	0.24089 (12)	0.50120 (7)	0.02242 (16)
H14A	-0.0465	0.2479	0.5664	0.034*
H14B	-0.1292	0.1227	0.4773	0.034*
H14C	-0.1735	0.2846	0.5220	0.034*
C15	-0.20773 (7)	0.41652 (14)	0.32699 (8)	0.02666 (19)
H15A	-0.1986	0.5387	0.3213	0.040*
H15B	-0.2666	0.3950	0.3538	0.040*
H15C	-0.2250	0.3646	0.2526	0.040*

supplementary materials

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0234 (3)	0.0354 (4)	0.0184 (3)	-0.0007 (2)	0.0091 (2)	0.0074 (2)
N1	0.0154 (3)	0.0231 (3)	0.0193 (3)	0.0014 (2)	0.0062 (2)	0.0022 (2)
C1	0.0176 (3)	0.0211 (3)	0.0172 (3)	-0.0011 (3)	0.0065 (3)	-0.0014 (3)
C2	0.0176 (3)	0.0274 (4)	0.0181 (3)	-0.0012 (3)	0.0049 (3)	0.0009 (3)
C3	0.0168 (3)	0.0271 (4)	0.0261 (4)	0.0013 (3)	0.0086 (3)	0.0050 (3)
C4	0.0208 (3)	0.0280 (4)	0.0262 (4)	0.0017 (3)	0.0133 (3)	0.0007 (3)
C5	0.0204 (3)	0.0254 (4)	0.0190 (3)	-0.0013 (3)	0.0102 (3)	-0.0012 (3)
C6	0.0162 (3)	0.0185 (3)	0.0166 (3)	-0.0019 (2)	0.0070 (2)	0.0003 (2)
C7	0.0183 (3)	0.0180 (3)	0.0156 (3)	-0.0021 (2)	0.0062 (3)	0.0002 (2)
C8	0.0162 (3)	0.0163 (3)	0.0143 (3)	-0.0005 (2)	0.0051 (2)	0.0006 (2)
C9	0.0195 (3)	0.0184 (3)	0.0152 (3)	0.0006 (2)	0.0052 (3)	0.0031 (2)
C10	0.0176 (3)	0.0193 (3)	0.0167 (3)	0.0026 (2)	0.0042 (3)	0.0029 (3)
C11	0.0154 (3)	0.0149 (3)	0.0148 (3)	-0.0003 (2)	0.0043 (2)	-0.0019 (2)
C12	0.0170 (3)	0.0169 (3)	0.0148 (3)	0.0006 (2)	0.0058 (2)	0.0016 (2)
C13	0.0158 (3)	0.0164 (3)	0.0154 (3)	0.0013 (2)	0.0049 (2)	0.0016 (2)
C14	0.0229 (3)	0.0252 (4)	0.0216 (4)	-0.0004 (3)	0.0112 (3)	0.0007 (3)
C15	0.0158 (3)	0.0350 (5)	0.0267 (4)	0.0032 (3)	0.0049 (3)	0.0048 (3)

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

O1—C7	1.2346 (10)	C8—C9	1.4031 (10)
N1—C11	1.3615 (10)	C8—C13	1.4066 (11)
N1—C14	1.4494 (11)	C9—C10	1.3781 (11)
N1—C15	1.4499 (11)	C9—H9	0.9300
C1—C2	1.3924 (11)	C10—C11	1.4163 (11)
C1—C6	1.3947 (11)	C10—H10	0.9300
C1—H1	0.9300	C11—C12	1.4165 (10)
C2—C3	1.3862 (12)	C12—C13	1.3816 (11)
C2—H2	0.9300	C12—H12	0.9300
C3—C4	1.3907 (13)	C13—H13	0.9300
C3—H3	0.9300	C14—H14A	0.9600
C4—C5	1.3868 (12)	C14—H14B	0.9600
C4—H4	0.9300	C14—H14C	0.9600
C5—C6	1.3965 (11)	C15—H15A	0.9600
C5—H5	0.9300	C15—H15B	0.9600
C6—C7	1.4976 (11)	C15—H15C	0.9600
C7—C8	1.4716 (11)		
C11—N1—C14	121.81 (7)	C10—C9—C8	122.12 (7)
C11—N1—C15	120.55 (7)	C10—C9—H9	118.9
C14—N1—C15	116.94 (7)	C8—C9—H9	118.9
C2—C1—C6	120.15 (8)	C9—C10—C11	120.77 (7)
C2—C1—H1	119.9	C9—C10—H10	119.6
C6—C1—H1	119.9	C11—C10—H10	119.6
C3—C2—C1	120.02 (8)	N1—C11—C10	120.86 (7)

C3—C2—H2	120.0	N1—C11—C12	121.89 (7)
C1—C2—H2	120.0	C10—C11—C12	117.25 (7)
C2—C3—C4	120.36 (8)	C13—C12—C11	121.14 (7)
C2—C3—H3	119.8	C13—C12—H12	119.4
C4—C3—H3	119.8	C11—C12—H12	119.4
C5—C4—C3	119.52 (8)	C12—C13—C8	121.49 (7)
C5—C4—H4	120.2	C12—C13—H13	119.3
C3—C4—H4	120.2	C8—C13—H13	119.3
C4—C5—C6	120.76 (8)	N1—C14—H14A	109.5
C4—C5—H5	119.6	N1—C14—H14B	109.5
C6—C5—H5	119.6	H14A—C14—H14B	109.5
C1—C6—C5	119.17 (7)	N1—C14—H14C	109.5
C1—C6—C7	123.28 (7)	H14A—C14—H14C	109.5
C5—C6—C7	117.47 (7)	H14B—C14—H14C	109.5
O1—C7—C8	120.51 (7)	N1—C15—H15A	109.5
O1—C7—C6	118.26 (7)	N1—C15—H15B	109.5
C8—C7—C6	121.18 (7)	H15A—C15—H15B	109.5
C9—C8—C13	117.23 (7)	N1—C15—H15C	109.5
C9—C8—C7	117.90 (7)	H15A—C15—H15C	109.5
C13—C8—C7	124.76 (7)	H15B—C15—H15C	109.5
C6—C1—C2—C3	-0.63 (13)	C6—C7—C8—C13	-12.54 (12)
C1—C2—C3—C4	1.18 (13)	C13—C8—C9—C10	-0.33 (12)
C2—C3—C4—C5	-0.29 (14)	C7—C8—C9—C10	175.94 (7)
C3—C4—C5—C6	-1.15 (13)	C8—C9—C10—C11	0.20 (12)
C2—C1—C6—C5	-0.79 (12)	C14—N1—C11—C10	177.99 (7)
C2—C1—C6—C7	-177.50 (7)	C15—N1—C11—C10	7.82 (12)
C4—C5—C6—C1	1.69 (12)	C14—N1—C11—C12	-1.90 (12)
C4—C5—C6—C7	178.59 (8)	C15—N1—C11—C12	-172.07 (8)
C1—C6—C7—O1	141.72 (9)	C9—C10—C11—N1	-179.62 (7)
C5—C6—C7—O1	-35.04 (11)	C9—C10—C11—C12	0.27 (11)
C1—C6—C7—C8	-40.56 (11)	N1—C11—C12—C13	179.28 (7)
C5—C6—C7—C8	142.68 (8)	C10—C11—C12—C13	-0.61 (11)
O1—C7—C8—C9	-10.84 (12)	C11—C12—C13—C8	0.49 (12)
C6—C7—C8—C9	171.49 (7)	C9—C8—C13—C12	-0.02 (11)
O1—C7—C8—C13	165.13 (8)	C7—C8—C13—C12	-176.00 (7)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C4—H4 \cdots O1 ⁱ	0.93	2.46	3.3730 (12)	168
C10—H10 \cdots Cg2 ⁱⁱ	0.93	2.98	3.6452 (9)	130

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

Fig. 1

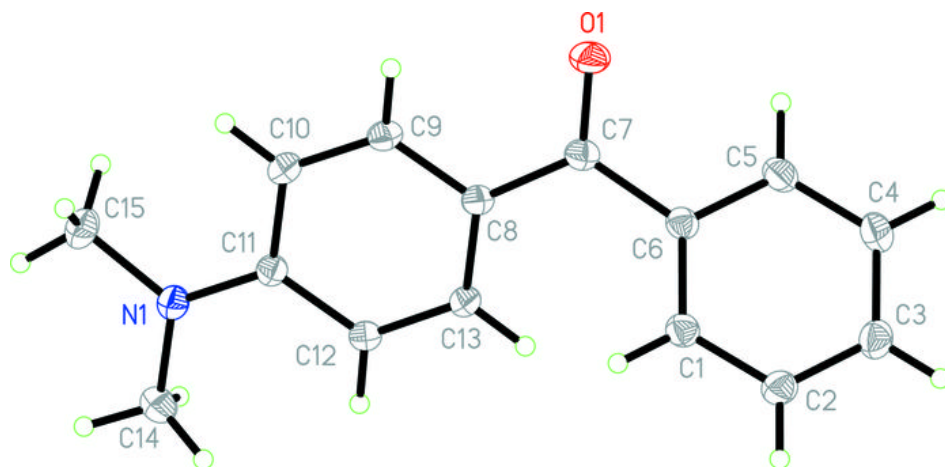


Fig. 2

