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(4-Fluorophenyl)(2-hydroxy-5-methylphenyl)methanone

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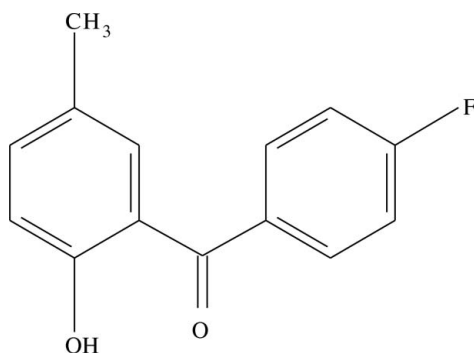
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.047; wR factor = 0.135; data-to-parameter ratio = 11.7.

In the title compound, $\text{C}_{14}\text{H}_{11}\text{FO}_2$, the dihedral angles between the central C_3O ketone residue and the fluoro- and hydroxy-substituted benzene rings are 50.44 (9) and 12.63 (10)°, respectively. The planes of the benzene rings subtend a dihedral angle of 58.88 (9)° and an intramolecular $\text{O}-\text{H}\cdots\text{O}$ hydrogen bond closes an $S(6)$ ring. No directional interactions beyond van der Waals packing contacts were identified in the crystal structure.

Related literature

For related structures, see: Dileep *et al.* (2013); Mahendra *et al.* (2005).



Experimental

Crystal data

 $\text{C}_{14}\text{H}_{11}\text{FO}_2$
 $M_r = 230.23$ Orthorhombic, $Pbca$
 $a = 5.9396$ (6) Å $b = 12.3808$ (15) Å
 $c = 30.522$ (3) Å
 $V = 2244.5$ (4) Å³
 $Z = 8$ Cu $K\alpha$ radiation
 $\mu = 0.85$ mm⁻¹
 $T = 296$ K
 $0.28 \times 0.25 \times 0.22$ mm

Data collection

Bruker X8 Proteum CCD
diffractometer
Absorption correction: multi-scan
(*SADABS*; Bruker, 2013)
 $T_{\min} = 0.798$, $T_{\max} = 0.836$ 8509 measured reflections
1822 independent reflections
1535 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.135$
 $S = 1.06$
1822 reflections156 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.18$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O1}-\text{H1}\cdots\text{O2}$	0.82	1.86	2.574 (2)	146

Data collection: *APEX2* (Bruker, 2013); cell refinement: *SAINT* (Bruker, 2013); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *Mercury* (Macrae *et al.*, 2008); software used to prepare material for publication: *Mercury*.

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Supporting information for this paper is available from the IUCr electronic archives (Reference: HB7191).

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supporting information

Acta Cryst. (2014). E70, o286 [doi:10.1107/S1600536814001883]

(4-Fluorophenyl)(2-hydroxy-5-methylphenyl)methanone

C. S. Dileep, V. Lakshmi Ranganatha, N. K. Lokanath, S. A. Khanum and M. A. Sridhar

S1. Comment

As part of our structural studies of benzophenone derivatives (Dileep *et al.*, 2013), the title compound was prepared and characterized by single-crystal X-ray diffraction.

The mean plane angle between the phenyl rings (/C1C2/C3/C4/C5/C6) and (C9/C10/C11/C12/C13/C14) is 58.88°.

The position of C8 atom is distorted trigonal planar geometry as indicated by bond angle values (O2—C8—C6)=121.05°, (O2—C8—C9)=118.37°, (N6—C8—C9)= 120.56°.

The conformation of the attachment of the two phenyl rings to the central carbonyl group can also be characterized by torsion angles (O2—C8—C6—C5) and (O2—C8—C9—C10) of -165.31° and -129.32°, respectively.

The crystal structure exhibits intramolecular O(1)—H(1)···O(2) hydrogen bonds. The bond angle between (O1—C1—C2) and (O1—C1—C6) is 118.61° and 122.22°. The bond length between (O1—C2) and (O2—C8) is 1.351 Å and 1.238 Å. The molecular packing when viewed down the *a* axis is shown in Fig. 2.

S2. Experimental

A mixture of anhydrous aluminium chloride (0.03 mol) and 4-fluoro-benzoic acid *p*-tolyl ester (0.02 mol) in dry nitrobenzene (40 ml) was protected from moisture by calcium chloride guard tube and refluxed for 45 min. At the end of this period the solution was cooled and decomposed by acidulated ice-cold water. Nitrobenzene was removed by steam distillation. The residual solid was crushed into powder, dissolved in ether and extracted with 10% sodium hydroxide. The basic aqueous solution was neutralized with 10% hydrochloric acid. The filtered solid was washed with distilled water and recrystallized from ethanol solution to afford pale yellow blocks of the title compound.

S3. Refinement

All H-atoms were positioned geometrically and refined using a riding model with C—H= 0.93–0.9600 Å and $U_{\text{iso}}(\text{H})=1.2U_{\text{eq}}(\text{C})$.

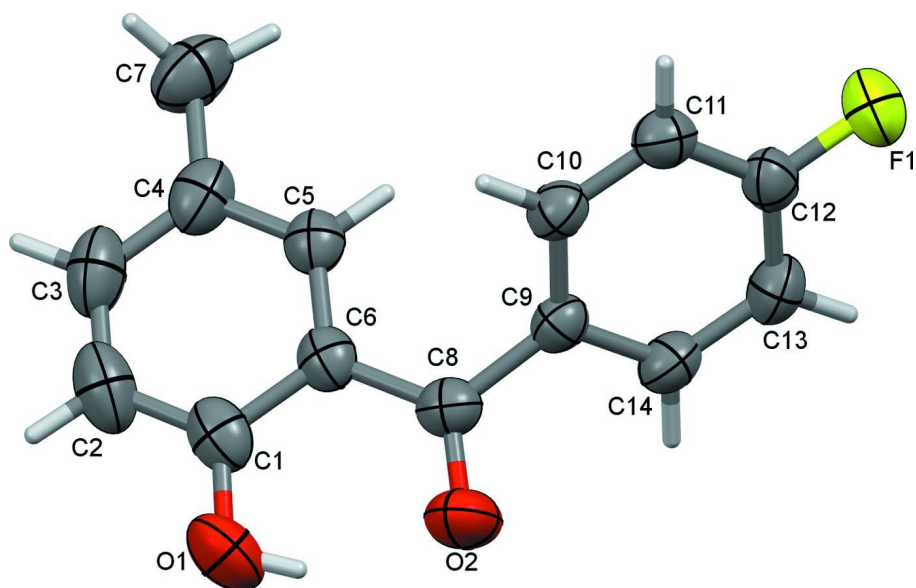


Figure 1

ORTEP view of the molecule with displacement ellipsoids drawn at the 50% probability level.

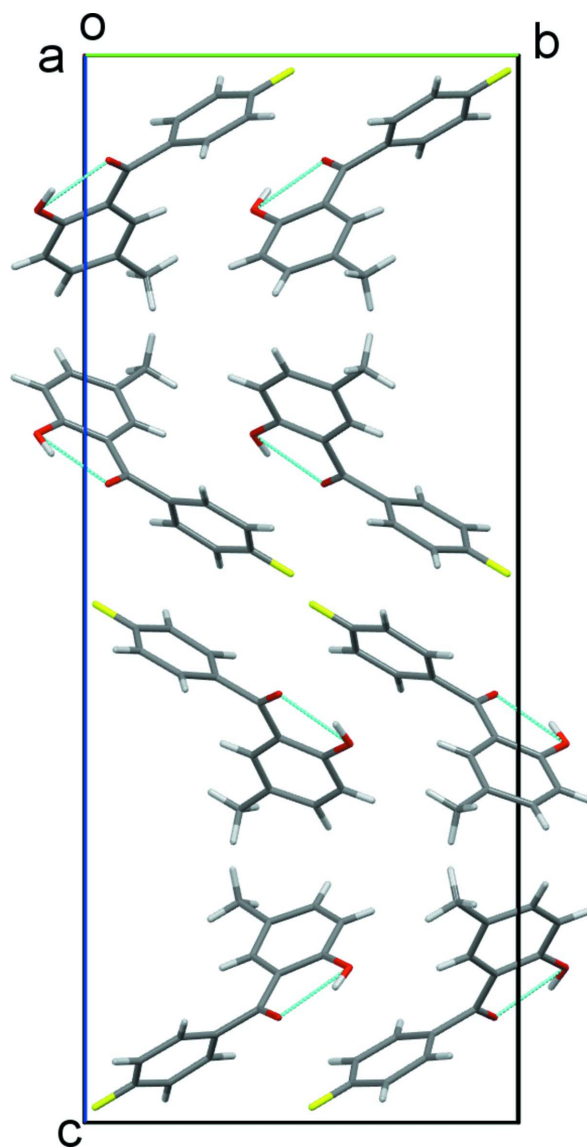


Figure 2

A molecular packing view of the title compound down the *a*-axis.

(4-Fluorophenyl)(2-hydroxy-5-methylphenyl)methanone

Crystal data

$C_{14}H_{11}FO_2$

$M_r = 230.23$

Orthorhombic, *Pbca*

Hall symbol: $-P\ 2ac\ 2ab$

$a = 5.9396\ (6)\ \text{\AA}$

$b = 12.3808\ (15)\ \text{\AA}$

$c = 30.522\ (3)\ \text{\AA}$

$V = 2244.5\ (4)\ \text{\AA}^3$

$Z = 8$

$F(000) = 960$

$D_x = 1.363\ \text{Mg m}^{-3}$

Cu $K\alpha$ radiation, $\lambda = 1.54178\ \text{\AA}$

Cell parameters from 8509 reflections

$\theta = 5.8\text{--}64.4^\circ$

$\mu = 0.85\ \text{mm}^{-1}$

$T = 296\ \text{K}$

Block, pale yellow

$0.28 \times 0.25 \times 0.22\ \text{mm}$

Data collection

Bruker X8 Proteum CCD
diffractometer

Radiation source: Bruker MicroStar microfocus
rotating anode

Helios multilayer optics monochromator

Detector resolution: 10.7 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2013)

$T_{\min} = 0.798$, $T_{\max} = 0.836$

8509 measured reflections

1822 independent reflections

1535 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.030$

$\theta_{\max} = 64.4^\circ$, $\theta_{\min} = 5.8^\circ$

$h = -6 \rightarrow 6$

$k = -14 \rightarrow 8$

$l = -34 \rightarrow 35$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.047$

$wR(F^2) = 0.135$

$S = 1.06$

1822 reflections

156 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

$w = 1/[\sigma^2(F_o^2) + (0.0857P)^2 + 0.3925P]$

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

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

$\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.21 \text{ e } \text{\AA}^{-3}$

Extinction correction: *SHELXL97* (Sheldrick,
2008), $FC^* = KFC[1 + 0.001XF^2\Lambda^3/\text{SIN}(2\Theta)]^{-1/4}$

Extinction coefficient: 0.0018 (4)

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement on F^2 for ALL reflections except those flagged by the user for potential systematic errors. Weighted R -factors wR and all goodnesses 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 observed criterion of $F^2 > \sigma(F^2)$ is used only for calculating $-R$ -factor-obs *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>	$U_{\text{iso}}^*/U_{\text{eq}}$
F1	0.2531 (2)	0.47588 (9)	0.01429 (4)	0.0737 (5)
O1	-0.0006 (3)	-0.11052 (11)	0.14523 (5)	0.0738 (6)
O2	-0.1159 (2)	0.05347 (12)	0.09849 (5)	0.0682 (5)
C1	0.1798 (4)	-0.05074 (14)	0.15707 (6)	0.0552 (6)
C2	0.3280 (4)	-0.09271 (16)	0.18766 (7)	0.0692 (8)
C3	0.5047 (4)	-0.0321 (2)	0.20256 (6)	0.0699 (8)
C4	0.5444 (3)	0.07352 (18)	0.18787 (6)	0.0576 (6)
C5	0.4015 (3)	0.11264 (14)	0.15595 (5)	0.0471 (6)
C6	0.2213 (3)	0.05270 (13)	0.13940 (5)	0.0453 (5)
C7	0.7297 (4)	0.1418 (2)	0.20654 (7)	0.0776 (9)
C8	0.0668 (3)	0.09745 (15)	0.10646 (6)	0.0469 (6)
C9	0.1246 (3)	0.19838 (13)	0.08253 (5)	0.0414 (5)
C10	0.3293 (3)	0.21172 (14)	0.06101 (5)	0.0455 (5)
C11	0.3715 (3)	0.30463 (15)	0.03721 (6)	0.0481 (6)

C12	0.2094 (3)	0.38360 (14)	0.03669 (5)	0.0482 (6)
C13	0.0073 (3)	0.37430 (16)	0.05775 (6)	0.0537 (6)
C14	-0.0356 (3)	0.27965 (15)	0.08026 (6)	0.0492 (6)
H1	-0.07480	-0.07740	0.12700	0.1110*
H2	0.30740	-0.16250	0.19810	0.0830*
H3	0.60200	-0.06200	0.22310	0.0840*
H5	0.42620	0.18170	0.14500	0.0560*
H7A	0.73900	0.20840	0.19050	0.1160*
H7B	0.87010	0.10380	0.20420	0.1160*
H7C	0.69850	0.15690	0.23680	0.1160*
H10	0.43820	0.15790	0.06260	0.0550*
H11	0.50590	0.31330	0.02200	0.0580*
H13	-0.09800	0.42990	0.05700	0.0640*
H14	-0.17390	0.27030	0.09410	0.0590*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
F1	0.0800 (9)	0.0591 (7)	0.0819 (8)	-0.0030 (6)	-0.0052 (6)	0.0248 (6)
O1	0.0938 (12)	0.0487 (8)	0.0789 (10)	-0.0156 (8)	0.0071 (8)	0.0023 (7)
O2	0.0575 (9)	0.0700 (9)	0.0770 (10)	-0.0208 (7)	-0.0081 (7)	0.0058 (7)
C1	0.0713 (13)	0.0429 (9)	0.0513 (10)	0.0047 (9)	0.0138 (9)	-0.0033 (8)
C2	0.1017 (18)	0.0495 (11)	0.0564 (11)	0.0150 (12)	0.0105 (12)	0.0081 (9)
C3	0.0838 (16)	0.0755 (15)	0.0504 (11)	0.0266 (13)	-0.0023 (10)	0.0123 (10)
C4	0.0560 (11)	0.0728 (12)	0.0440 (9)	0.0156 (10)	0.0005 (8)	-0.0009 (9)
C5	0.0489 (10)	0.0486 (10)	0.0437 (9)	0.0066 (8)	0.0050 (7)	0.0007 (7)
C6	0.0500 (10)	0.0418 (9)	0.0441 (9)	0.0054 (8)	0.0074 (7)	-0.0017 (7)
C7	0.0625 (13)	0.1073 (19)	0.0630 (13)	0.0040 (13)	-0.0148 (10)	-0.0016 (12)
C8	0.0427 (10)	0.0482 (10)	0.0499 (10)	-0.0023 (8)	0.0017 (7)	-0.0052 (7)
C9	0.0385 (9)	0.0446 (9)	0.0410 (8)	0.0014 (7)	-0.0049 (6)	-0.0032 (6)
C10	0.0397 (9)	0.0460 (9)	0.0507 (9)	0.0049 (7)	-0.0016 (7)	-0.0032 (7)
C11	0.0431 (10)	0.0555 (10)	0.0458 (9)	-0.0032 (8)	0.0002 (7)	-0.0001 (7)
C12	0.0533 (11)	0.0448 (9)	0.0464 (9)	-0.0029 (8)	-0.0095 (8)	0.0055 (7)
C13	0.0511 (11)	0.0516 (10)	0.0584 (11)	0.0127 (9)	-0.0062 (8)	0.0022 (8)
C14	0.0375 (9)	0.0591 (11)	0.0510 (10)	0.0048 (8)	-0.0015 (7)	-0.0009 (8)

Geometric parameters (Å, °)

F1—C12	1.357 (2)	C10—C11	1.383 (3)
O1—C1	1.352 (3)	C11—C12	1.372 (3)
O2—C8	1.238 (2)	C12—C13	1.367 (2)
O1—H1	0.8200	C13—C14	1.382 (3)
C1—C6	1.411 (2)	C2—H2	0.9300
C1—C2	1.384 (3)	C3—H3	0.9300
C2—C3	1.368 (3)	C5—H5	0.9300
C3—C4	1.402 (3)	C7—H7A	0.9600
C4—C5	1.380 (2)	C7—H7B	0.9600
C4—C7	1.500 (3)	C7—H7C	0.9600

C5—C6	1.397 (2)	C10—H10	0.9300
C6—C8	1.470 (2)	C11—H11	0.9300
C8—C9	1.488 (2)	C13—H13	0.9300
C9—C14	1.387 (2)	C14—H14	0.9300
C9—C10	1.392 (2)		
C1—O1—H1	110.00	C11—C12—C13	123.43 (17)
O1—C1—C6	122.23 (18)	C12—C13—C14	117.85 (17)
C2—C1—C6	119.16 (19)	C9—C14—C13	120.90 (16)
O1—C1—C2	118.61 (17)	C1—C2—H2	120.00
C1—C2—C3	120.41 (19)	C3—C2—H2	120.00
C2—C3—C4	122.29 (19)	C2—C3—H3	119.00
C3—C4—C5	116.72 (18)	C4—C3—H3	119.00
C3—C4—C7	121.83 (18)	C4—C5—H5	119.00
C5—C4—C7	121.44 (19)	C6—C5—H5	119.00
C4—C5—C6	122.69 (17)	C4—C7—H7A	109.00
C1—C6—C5	118.54 (16)	C4—C7—H7B	109.00
C5—C6—C8	121.70 (15)	C4—C7—H7C	109.00
C1—C6—C8	119.63 (16)	H7A—C7—H7B	109.00
O2—C8—C9	118.37 (16)	H7A—C7—H7C	110.00
C6—C8—C9	120.56 (15)	H7B—C7—H7C	110.00
O2—C8—C6	121.05 (17)	C9—C10—H10	120.00
C8—C9—C14	118.41 (16)	C11—C10—H10	120.00
C10—C9—C14	119.33 (15)	C10—C11—H11	121.00
C8—C9—C10	122.21 (15)	C12—C11—H11	121.00
C9—C10—C11	120.32 (16)	C12—C13—H13	121.00
C10—C11—C12	118.12 (16)	C14—C13—H13	121.00
F1—C12—C11	118.14 (15)	C9—C14—H14	120.00
F1—C12—C13	118.43 (16)	C13—C14—H14	120.00
O1—C1—C2—C3	-175.88 (19)	C5—C6—C8—C9	13.4 (3)
C6—C1—C2—C3	4.0 (3)	O2—C8—C9—C10	-129.33 (19)
O1—C1—C6—C5	174.98 (17)	O2—C8—C9—C14	48.0 (2)
O1—C1—C6—C8	-1.0 (3)	C6—C8—C9—C10	52.0 (2)
C2—C1—C6—C5	-4.9 (3)	C6—C8—C9—C14	-130.74 (18)
C2—C1—C6—C8	179.17 (18)	C8—C9—C10—C11	176.58 (16)
C1—C2—C3—C4	-0.1 (3)	C14—C9—C10—C11	-0.7 (2)
C2—C3—C4—C5	-2.7 (3)	C8—C9—C14—C13	-178.76 (17)
C2—C3—C4—C7	176.0 (2)	C10—C9—C14—C13	-1.4 (3)
C3—C4—C5—C6	1.7 (3)	C9—C10—C11—C12	2.0 (3)
C7—C4—C5—C6	-177.01 (17)	C10—C11—C12—F1	178.33 (15)
C4—C5—C6—C1	2.0 (3)	C10—C11—C12—C13	-1.3 (3)
C4—C5—C6—C8	177.92 (17)	F1—C12—C13—C14	179.67 (16)
C1—C6—C8—O2	10.5 (3)	C11—C12—C13—C14	-0.7 (3)
C1—C6—C8—C9	-170.79 (16)	C12—C13—C14—C9	2.1 (3)
C5—C6—C8—O2	-165.31 (17)		

Hydrogen-bond geometry (Å, °)

<i>D—H···A</i>	<i>D—H</i>	<i>H···A</i>	<i>D···A</i>	<i>D—H···A</i>
O1—H1···O2	0.82	1.86	2.574 (2)	146