Acta Crystallographica Section E **Structure Reports** Online

ISSN 1600-5368

(4-Hydroxy-2,5-dimethylphenyl)phenylmethanone

Rodolfo Moreno-Fuquen,^a* Leidy J. Valencia,^a Alan R. Kennedy,^b Denise Gilmore^b and R. H. De Almeida Santos^c

^aDepartamento de Química – Facultad de Ciencias, Universidad del Valle, Apartado 25360, Santiago de Cali, Colombia, ^bWestCHEM, Department of Pure and Applied Chemistry, University of Strathclyde, 295 Cathedral Street, Glasgow G1 1XL, Scotland, and ^cInstituto de Química de São Carlos, Universidade de São Paulo, USP, São Carlos, SP, Brazil

Correspondence e-mail: rodimo26@yahoo.es

Received 16 September 2009; accepted 28 September 2009

Key indicators: single-crystal X-ray study; T = 123 K; mean σ (C–C) = 0.002 Å; R factor = 0.032; wR factor = 0.058; data-to-parameter ratio = 13.0.

The title compound, C₁₅H₁₄O₂, was obtained by Friedel-Crafts acylation between 2,5-dimethylphenol and benzoyl chloride in the presence of aluminium chloride as a catalyst. The dihedral angle between the benzene rings is $61.95 (4)^{\circ}$. In the crystal, $O-H \cdots O$ hydrogen bonding and $C-H \cdots O$ weak interactions lead to polymeric C(6), C(8) and C(11) chains along the *a*, *b* and *c*-axis directions, respectively.

Related literature

For background information on the anti-fungal and antiinflamatory biological activity of benzophenones, see: Naldoni et al. (2009); Selvi et al. (2003); Naveen et al. (2006). For 104 benzophenone molecules, see: Cox et al. (2008). For hydrogenbond motifs, see: Etter (1990).



Experimental

Crystal data

C15H14O2 $M_r = 226.26$ Orthorhombic, Pbca a = 12.1392 (10) Åb = 8.1386 (7) Å c = 23.665 (2) Å

V = 2338.0 (3) Å³ Z = 8Mo $K\alpha$ radiation $\mu = 0.08 \text{ mm}^{-1}$ T = 123 K $0.25\,\times\,0.12\,\times\,0.05$ mm



Data collection

Oxford Diffraction Gemini S
diffractometer
Absorption correction: multi-scan
(CrysAlis CCD; Oxford
Diffraction, 2009)
$T_{\min} = 0.904, T_{\max} = 1.000$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$	158 parameters
$wR(F^2) = 0.058$	H-atom parameters constrained
S = 0.73	$\Delta \rho_{\rm max} = 0.15 \text{ e} \text{ Å}^{-3}$
2059 reflections	$\Delta \rho_{\rm min} = -0.14 \text{ e } \text{\AA}^{-3}$

9067 measured reflections

 $R_{\rm int} = 0.061$

2059 independent reflections 1061 reflections with $I > 2\sigma(I)$

Table 1

Hydrogen-bond geometry (Å, °).

			DI	$D = 11 \cdots 11$
$O2-H2\cdots O1^i$	0.84	1.92	2.6973 (15)	154
$C15-H15B\cdots O1^{ii}$	0.98	2.62	3.352 (2)	132
$C4-H4\cdots O2^{iii}$	0.95	2.67	3.454 (2)	140

Data collection: CrysAlis CCD (Oxford Diffraction, 2009); cell refinement: CrysAlis CCD; data reduction: CrysAlis RED (Oxford Diffraction, 2009); program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: ORTEP-3 for Windows

(Farrugia, 1997) and Mercury (Macrae et al., 2006); software used to

prepare material for publication: PARST95 (Nardelli, 1995).

RMF is grateful to the Spanish Research Council (CSIC) for the use of a free-of-charge licence to the Cambridge Structural Database (Allen, 2002). RMF also thanks the Universidad del Valle, Colombia, and the Instituto de Química de São Carlos, Brazil, for partial financial support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HG2568).

References

Allen, F. H. (2002). Acta Cryst. B58, 380-388.

Cox, P. J., Kechagias, D. & Kelly, O. (2008). Acta Cryst. B64, 206-216.

- Etter, M. (1990). Acc. Chem. Res. 23, 120-126.
- Farrugia, L. J. (1997). J. Appl. Cryst. 30, 565.
- Macrae, C. F., Edgington, P. R., McCabe, P., Pidcock, E., Shields, G. P., Taylor, R., Towler, M. & van de Streek, J. (2006). J. Appl. Cryst. 39, 453-457.

Naldoni, F. J., Claudino, A. L. R., Cruz, J. W., Chavasco, J. K., Faria e Silva, P. M., Veloso, M. P. & Dos Santos, M. H. (2009). J. Med. Food, 12, 403-407. Nardelli, M. (1995). J. Appl. Cryst. 28, 659.

Naveen, S., Khanum, S. A., Devaiah, V. T., Shashikanth, S., Anandalwar, S. M. & Prasad, S. (2006). Anal. Sci. 22, 183-184.

Oxford Diffraction (2009). CrysAlis CCD and CrysAlis RED. Oxford Diffraction Ltd, Yarnton, England,

Selvi, A. T., Joseph, G. S. & Jayaprakasha, G. K. (2003). Food Microbiol. 20, 455-460.

Sheldrick, G. M. (2008). Acta Cryst. A64, 112-122.

supporting information

Acta Cryst. (2009). E65, o2614 [https://doi.org/10.1107/S1600536809039488]

(4-Hydroxy-2,5-dimethylphenyl)phenylmethanone

Rodolfo Moreno-Fuquen, Leidy J. Valencia, Alan R. Kennedy, Denise Gilmour and R. H. De Almeida Santos

S1. Comment

The title compound, $C_{15}H_{14}O_2$, (4-Hydroxy-2,5-dimethyl-phenyl)-phenyl-methanone), (I), is part of a series of studies on benzophenone, which have been made in our research group. Benzophenone analogue systems show various anti-fungal and anti-inflamatory biological activities (Naldoni et al., 2009 and Selvi et al., 2003). The presence of various substituents in the benzophenone nucleus is essential to determining the quantitative structure-activity relationships of these systems. Some studies were carried out to show that methyl-substituted benzophenones exhibit anti-fungal properties (Naveen et al., 2006). In order to present the molecular conformation of (I), to analyse the type of hydrogenbonds formed in (I) and to study its supramolecular behavior, the title compound was synthesized. The molecular structure of the title compound is shown in Fig. 1. The bond lengths and bond angles of (I) are in good agreement with the standard values and correspond to those observed in (4-Hydroxy-3-methylphenyl)(4- methylphenyl)methanone (Naveen et al., 2006). The two aromatic rings in the title structure form a dihedral angle of 61.95 (4)°. This value is greater than the value presented in the stable, orthorhombic form of unsubstituted benzophenone (54°) and follows the standard behavior of the majority of benzophenone molecules [104 benzophenone molecules, Cox et al., 2008]. The title molecule is characterized by the formation of O—H···O hydrogen-bonds and other C—H···O weak interactions (Table 1, Nardelli, 1995). The strongest hydrogen bond O—H…O interaction is responsible for crystal growth in [100] direction. Indeed, in a first substructure, atom O2 in the molecule at (x, y, z) acts as hydrogen bond donor to carbonyl O1 atom in the molecule at (x - 1/2, -y + 1/2, -z + 1). The propagation of this interaction forms a C(8) (Etter, 1990) chain running along [100] direction (Fig. 2). In a second substructure, atom C15 in the molecule at (x, y, z) links with weak interaction to carbonyl O1 atom in the molecule at (-x + 3/2, y - 1/2, z). The propagation of this interaction forms C(6) continuous chains via C15—H15B···O1 and running along [010] direction (Fig. 3). Finally in a third sub-structure, atom C4 in the molecule at (x, y, z) links with weak interaction to hydroxyl O2 atom in the molecule at (x, -y + 3/2, z - 1/2). The propagation of this interaction forms C(11) continuous chains and running along [001] direction. All of these interactions in [100], [010] and [001] directions define the bulk structure of the crystal.

S2. Experimental

2,5-dimethylphenol (0.50 g, 4.10 mmol) was added to a solution of anhydrous aluminium chloride (0.40 g, 3.00 mmol) in dry dichloromethane (25 ml). The resulting solution was cooled and then a benzoyl chloride (0.80 g, 5.70 mmol) was slowly added at $0-5^{\circ}$ C. After complete addition, the mixture was allowed to stir at room temperature for 0.5 h, and then it was heated up to 50° C for 1 h. The reaction mixture was poured onto ice (100 g) and conc. HCl (10 ml). The crude product was isolated by extraction with dichloromethane. The combined organic layers were washed with 10% aqueous NaOH, water, and then the solution was dried over Na₂SO₄ and it was evaporated at room temperature.

S3. Refinement

All H-atoms were located from difference maps and were positioned geometrically and refined using a riding model with C–H= 0.93–0.97 Å and $U_{iso}(H)$ = 1.2 $U_{eq}(C)$.



Figure 1

An *ORTEP-3* (Farrugia, 1997) plot of the title (I) compound, with the atomic labelling scheme. The shapes of the ellipsoids correspond to 50% probability contours of atomic displacement and, for the sake of clarity, H atoms are shown as spheres of arbitrary radius.



Figure 2

Part of the crystal structure of (I), showing the formation of C(8) chains running along [100] direction. Symmetry code: (i) x + 1/2, -y + 1/2, -z + 1; (ii) x - 1/2, -y + 1/2, -z + 1



Figure 3

Part of the crystal structure of (I), showing the formation of C(6) chain running along [010]. Symmetry code: (i) -x + 3/2, y - 1/2, z; (ii) -x + 3/2, y + 1/2, z.



Figure 4

Part of the crystal structure of (I), showing the formation of C(11) chain running along [001]. Symmetry code: (i) x, -y + 3/2, z - 1/2; (ii) x, -y + 3/2, z + 1/2

(4-Hydroxy-2,5-dimethylphenyl)phenylmethanone

Crystal data $C_{15}H_{14}O_2$ $D_{\rm x} = 1.286 {\rm Mg} {\rm m}^{-3}$ $M_r = 226.26$ Melting point: 443.0(10) K Orthorhombic, Pbca Mo *K* α radiation, $\lambda = 0.71073$ Å Hall symbol: -P 2ac 2ab Cell parameters from 1709 reflections $\theta = 2.5 - 30.7^{\circ}$ *a* = 12.1392 (10) Å *b* = 8.1386 (7) Å $\mu = 0.08 \text{ mm}^{-1}$ T = 123 Kc = 23.665 (2) Å $V = 2338.0(3) \text{ Å}^3$ Shard, colourless Z = 8 $0.25 \times 0.12 \times 0.05 \text{ mm}$ F(000) = 960

Data collection

Oxford Diffraction Gemini S diffractometer Radiation source: fine-focus sealed tube Graphite monochromator ω scans Absorption correction: multi-scan (<i>CrysAlis CCD</i> ; Oxford Diffraction, 2009) $T_{\min} = 0.904, T_{\max} = 1.000$ Refinement	9067 measured reflections 2059 independent reflections 1061 reflections with $I > 2\sigma(I)$ $R_{int} = 0.061$ $\theta_{max} = 25.0^{\circ}, \ \theta_{min} = 3.1^{\circ}$ $h = -12 \rightarrow 14$ $k = -9 \rightarrow 9$ $l = -28 \rightarrow 26$
Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.032$ $wR(F^2) = 0.058$ S = 0.73 2059 reflections 158 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 $w = 1/[\sigma^2(F_o^2) + (0.0224P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.15$ e Å ⁻³ $\Delta\rho_{min} = -0.14$ e Å ⁻³ Extinction correction: <i>SHELXL97</i> (Sheldrick, 2008), Fc*=kFc[1+0.001xFc^2\lambda^3/sin(2\theta)]^{-1/4} Extinction coefficient: 0.0019 (2)

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 $(Å^2)$

	x	у	Ζ	$U_{ m iso}$ */ $U_{ m eq}$	
01	0.83004 (8)	0.77322 (14)	0.37654 (5)	0.0301 (3)	
O2	0.51698 (9)	0.86337 (14)	0.58360 (5)	0.0261 (3)	
H2	0.4559	0.8151	0.5849	0.039*	
C1	0.67652 (14)	0.8006 (2)	0.31719 (7)	0.0211 (4)	
C2	0.73020 (14)	0.7315 (2)	0.27099 (7)	0.0291 (5)	
H2A	0.7986	0.6768	0.2761	0.035*	
C3	0.68450 (16)	0.7422 (2)	0.21789 (7)	0.0370 (5)	
H3	0.7213	0.6946	0.1865	0.044*	
C4	0.58475 (16)	0.8226 (2)	0.21010 (8)	0.0363 (5)	
H4	0.5527	0.8278	0.1735	0.044*	
C5	0.53206 (15)	0.8948 (2)	0.25549 (7)	0.0306 (5)	
H5	0.4649	0.9524	0.2499	0.037*	
C6	0.57691 (14)	0.8834 (2)	0.30907 (7)	0.0248 (5)	
H6	0.5400	0.9317	0.3403	0.030*	
C7	0.72879 (13)	0.78937 (19)	0.37381 (7)	0.0210 (4)	

C8	0.66409 (13)	0.80471 (19)	0.42695 (7)	0.0186 (4)	
C9	0.71040 (13)	0.9034 (2)	0.46920 (7)	0.0209 (4)	
H9	0.7782	0.9572	0.4616	0.025*	
C10	0.66178 (13)	0.9261 (2)	0.52152 (7)	0.0185 (4)	
C11	0.56262 (13)	0.8428 (2)	0.53126 (7)	0.0195 (4)	
C12	0.51675 (13)	0.74203 (19)	0.49063 (7)	0.0210 (4)	
H12	0.4505	0.6849	0.4990	0.025*	
C13	0.56518 (13)	0.7218 (2)	0.43752 (7)	0.0189 (4)	
C14	0.71190 (13)	1.0335 (2)	0.56630(7)	0.0270 (5)	
H14A	0.7342	0.9656	0.5986	0.040*	
H14B	0.6577	1.1150	0.5787	0.040*	
H14C	0.7766	1.0898	0.5509	0.040*	
C15	0.51105 (14)	0.6050(2)	0.39635 (7)	0.0266 (5)	
H15A	0.4730	0.5180	0.4173	0.040*	
H15B	0.5673	0.5560	0.3719	0.040*	
H15C	0.4577	0.6653	0.3732	0.040*	

Atomic displacement parameters (\mathring{A}^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
01	0.0175 (6)	0.0449 (8)	0.0279 (7)	0.0032 (7)	-0.0010 (6)	0.0015 (7)
O2	0.0221 (7)	0.0330 (8)	0.0231 (7)	-0.0020 (6)	0.0038 (6)	0.0001 (6)
C1	0.0213 (10)	0.0219 (10)	0.0200 (10)	-0.0005 (9)	-0.0006 (9)	0.0020 (9)
C2	0.0321 (10)	0.0282 (12)	0.0269 (11)	0.0059 (9)	0.0025 (10)	0.0014 (9)
C3	0.0531 (13)	0.0374 (13)	0.0206 (11)	0.0084 (11)	0.0015 (11)	-0.0035 (9)
C4	0.0521 (14)	0.0336 (13)	0.0231 (12)	0.0027 (11)	-0.0110 (11)	0.0034 (10)
C5	0.0343 (12)	0.0256 (12)	0.0318 (12)	0.0013 (9)	-0.0102 (10)	0.0043 (10)
C6	0.0256 (10)	0.0246 (11)	0.0241 (11)	-0.0005 (9)	0.0002 (10)	-0.0013 (9)
C7	0.0231 (9)	0.0179 (10)	0.0219 (10)	0.0001 (8)	-0.0003 (9)	0.0007 (9)
C8	0.0176 (9)	0.0187 (10)	0.0196 (10)	0.0029 (8)	-0.0009 (9)	0.0019 (9)
С9	0.0172 (9)	0.0203 (10)	0.0253 (11)	-0.0006 (8)	-0.0015 (9)	0.0069 (9)
C10	0.0199 (10)	0.0173 (10)	0.0185 (10)	0.0035 (8)	-0.0030 (9)	0.0021 (8)
C11	0.0204 (10)	0.0207 (10)	0.0173 (10)	0.0063 (8)	0.0026 (9)	0.0037 (9)
C12	0.0170 (9)	0.0206 (11)	0.0253 (10)	-0.0004 (9)	0.0004 (8)	0.0049 (9)
C13	0.0173 (9)	0.0181 (10)	0.0215 (10)	0.0023 (8)	-0.0029 (8)	0.0006 (8)
C14	0.0252 (10)	0.0274 (11)	0.0283 (11)	-0.0007 (9)	0.0004 (9)	0.0027 (9)
C15	0.0267 (10)	0.0248 (11)	0.0283 (11)	-0.0032 (9)	-0.0006 (9)	-0.0002 (9)

Geometric parameters (Å, °)

01—C7	1.2377 (17)	C8—C13	1.400 (2)	
O2—C11	1.3674 (18)	C8—C9	1.400 (2)	
O2—H2	0.8400	C9—C10	1.384 (2)	
C1—C2	1.391 (2)	С9—Н9	0.9500	
C1—C6	1.397 (2)	C10—C11	1.401 (2)	
C1—C7	1.486 (2)	C10—C14	1.503 (2)	
С2—С3	1.376 (2)	C11—C12	1.381 (2)	
C2—H2A	0.9500	C12—C13	1.397 (2)	

supporting information

C3—C4	1.388 (2)	С12—Н12	0.9500
C3—H3	0.9500	C13—C15	1.511 (2)
C4—C5	1 381 (2)	C14—H14A	0.9800
C4—H4	0.9500	C14—H14B	0.9800
C5—C6	1.383 (2)	C14—H14C	0.9800
C5—H5	0.9500	C15—H15A	0.9800
C6—H6	0.9500	C15—H15B	0.9800
C7—C8	1 488 (2)	C15—H15C	0.9800
			012000
С11—О2—Н2	109.5	С10—С9—Н9	118.5
$C_{2}-C_{1}-C_{6}$	119.48 (15)	С8—С9—Н9	118.5
C2-C1-C7	118.93 (15)	C9-C10-C11	116.70 (16)
C6—C1—C7	121.56 (15)	C9-C10-C14	122.40 (15)
C_{3} $-C_{2}$ $-C_{1}$	120.20 (16)	C11—C10—C14	120.90(15)
C3-C2-H2A	119.9	02-C11-C12	122.69(15)
C1 - C2 - H2A	119.9	02 - C11 - C10	115.98 (15)
$C^2 - C^3 - C^4$	120 17 (17)	C12 - C11 - C10	121 29 (16)
C2—C3—H3	119.9	$C_{11} - C_{12} - C_{13}$	121.29 (16)
C4-C3-H3	119.9	C11 - C12 - H12	119.1
C_{5} C_{4} C_{3}	120.07 (17)	C13 - C12 - H12	119.1
$C_{2} = C_{4} = C_{2}$	120.07 (17)	C_{12} C_{12} C_{13} C_{8}	117.69 (15)
$C_3 - C_4 - H_4$	120.0	$C_{12} = C_{13} = C_{15}$	118 11 (15)
C4 - C5 - C6	120.0	C8-C13-C15	124 11 (14)
C4 - C5 - H5	110.0	C_{10} C_{14} H_{14A}	109 5
C6 C5 H5	119.9	C10 C14 H14B	109.5
C_{0}	119.9	H_{14} C_{14} H_{14} H_{14}	109.5
C_{5}	119.95 (10)	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	109.5
C_{1} C_{6} H_{6}	120.0		109.5
$C_1 = C_0 = H_0$	120.0	H14A - C14 - H14C	109.5
01 - 07 - 01	110.34(10) 110.27(16)	114D - C14 - 114C	109.5
01 - 07 - 08	119.27(10) 122.12(14)	С12 С15 Ц15Р	109.5
$C1 = C^{2} = C^{2}$	122.12(14)		109.5
C13 - C8 - C7	119.30(13) 124.28(15)	HI3A—CI3—HI3B	109.5
$C_{13} = C_{8} = C_{7}$	124.28(15)	LIS-CIS-HISC	109.5
$C_{9} = C_{8} = C_{7}$	110.08 (15)	HISA—CIS—HISC	109.5
C10-C9-C8	122.95 (16)	HISB-CIS-HISC	109.5
C6 C1 C2 C3	-11(3)	C13 C8 C9 C10	-12(2)
$C_0 = C_1 = C_2 = C_3$	-170.02(16)	$C_{13} = C_{3} = C_{10} = C_{10}$	1.2(2)
$C_1 = C_2 = C_3$	1/9.02(10)	$C_{1}^{2} = C_{1}^{2} = C_{1$	1/8.13(13)
$C_1 = C_2 = C_3 = C_4$	0.2(3)	$C_{8} = C_{9} = C_{10} = C_{14}$	-170.48(16)
$C_2 = C_3 = C_4 = C_5$	1.3(3) -1.8(3)	$C_{0} = C_{10} = C_{11} = C_{14}$	-1/9.46(10) 178 50(14)
$C_{3} - C_{4} - C_{5} - C_{6}$	-1.8(3)	C_{9} C_{10} C_{11} C_{2} C_{14} C_{10} C_{11} C_{2}	1/8.30(14)
C4 - C3 - C0 - C1	0.9(3)	$C_{14} = C_{10} = C_{11} = C_{12}$	-1.2(2)
$\begin{array}{cccccccccccccccccccccccccccccccccccc$	0.3(3)	$C_{14} = C_{10} = C_{11} = C_{12}$	0.0(2)
$C_1 = C_1 = C_2 = C_1$	1/0.40(10)	$C_{14} = C_{10} = C_{11} = C_{12}$	-1/9.08(13)
$C_{2} - C_{1} - C_{1} - C_{1}$	24.8(2) -152(00(16)	02-011-012-013	-1/9.51(14) -1.8(2)
$C_0 - C_1 - C_7 - C_1^0$	-133.09 (16)	C_{10} $-C_{11}$ $-C_{12}$ $-C_{13}$ C_{12} C_{12} C_{13}	-1.8(2)
$U_2 - U_1 - U_1 - U_8$	-158.14(16)	C11 - C12 - C13 - C8	1.4 (2)
Co-CI-C/-C8	24.0 (2)	C11—C12—C13—C15	1/8.17 (14)

supporting information

O1—C7—C8—C13	-135.69 (17)	C9—C8—C13—C12	0.0 (2)
C1—C7—C8—C13	47.3 (2)	C7—C8—C13—C12	176.75 (15)
O1—C7—C8—C9	41.1 (2)	C9—C8—C13—C15	-176.54 (15)
C1—C7—C8—C9	-135.91 (16)	C7—C8—C13—C15	0.2 (2)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D····A	<i>D</i> —H··· <i>A</i>
O2—H2···O1 ⁱ	0.84	1.92	2.6973 (15)	154
C15—H15 <i>B</i> ····O1 ⁱⁱ	0.98	2.62	3.352 (2)	132
C4—H4····O2 ⁱⁱⁱ	0.95	2.67	3.454 (2)	140

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