

Crystal structure of (*E*)-*N'*-benzylidene-2-methoxybenzohydrazideMuhammad Taha,^{a,b} M. Syukri Baharudin,^a Hamidah Mohd Zaki,^{a,b} Bohari M. Yamin^c and Humera Naz^{a,d,*}

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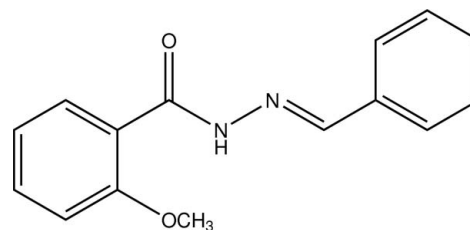
In the title benzoylhydrazide derivative, C₁₅H₁₄N₂O₂, the dihedral angle between the planes of the two phenyl rings is 12.56 (9)°. The azomethine double bond adopts an *E* configuration stabilized by an N—H···O hydrogen bond. In the crystal, the components are linked by C—H···O interactions to form chains along the *b* axis.

Keywords: crystal structure; benzohydrazide; Schiff base; hydrogen bonding.

CCDC reference: 1020647

1. Related literature

For applications and biological activities of Schiff bases, see: Taha *et al.* (2013, 2014); Musharraf *et al.* (2012); Kaymakcioglu *et al.* (2006); Kucukguzel *et al.* (2003, 2004); Melnyk *et al.* (2006); Pandeya *et al.* (1999); Tarafder *et al.* (2002); Terzioglu & Gursoy (2003); Todeschini *et al.* (1998). For the crystal structures of related compounds, see: Taha *et al.* (2013).



2. Experimental

2.1. Crystal data

C ₁₅ H ₁₄ N ₂ O ₂	<i>V</i> = 2655.2 (3) Å ³
<i>M_r</i> = 254.28	<i>Z</i> = 8
Orthorhombic, <i>Pbca</i>	Mo <i>K</i> α radiation
<i>a</i> = 13.3135 (9) Å	<i>μ</i> = 0.09 mm ⁻¹
<i>b</i> = 9.9581 (6) Å	<i>T</i> = 296 K
<i>c</i> = 20.0278 (14) Å	0.50 × 0.48 × 0.31 mm

2.2. Data collection

Bruker SMART APEX CCD area-detector diffractometer	38941 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2000)	2462 independent reflections
<i>T_{min}</i> = 0.94, <i>T_{max}</i> = 0.97	1819 reflections with <i>I</i> > 2σ(<i>I</i>)
	<i>R_{int}</i> = 0.049

2.3. Refinement

<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)] = 0.038	174 parameters
<i>wR</i> (<i>F</i> ²) = 0.103	H-atom parameters constrained
<i>S</i> = 1.08	Δ <i>ρ</i> _{max} = 0.13 e Å ⁻³
2462 reflections	Δ <i>ρ</i> _{min} = -0.14 e Å ⁻³

Table 1

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>
N2—H2A···O2	0.86	1.96	2.6278 (17)	134
C7—H7A···O1 ⁱ	0.93	2.44	3.1690 (19)	135

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

Data collection: *SMART* (Bruker, 2000); cell refinement: *SAINTE* (Bruker, 2000); data reduction: *SAINTE*; 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*, *PARST* (Nardelli, 1995) and *PLATON* (Spek, 2009).

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Supporting information for this paper is available from the IUCr electronic archives (Reference: BG2535).

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Crystal structure of (*E*)-*N'*-benzylidene-2-methoxybenzohydrazide

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S1. Comment

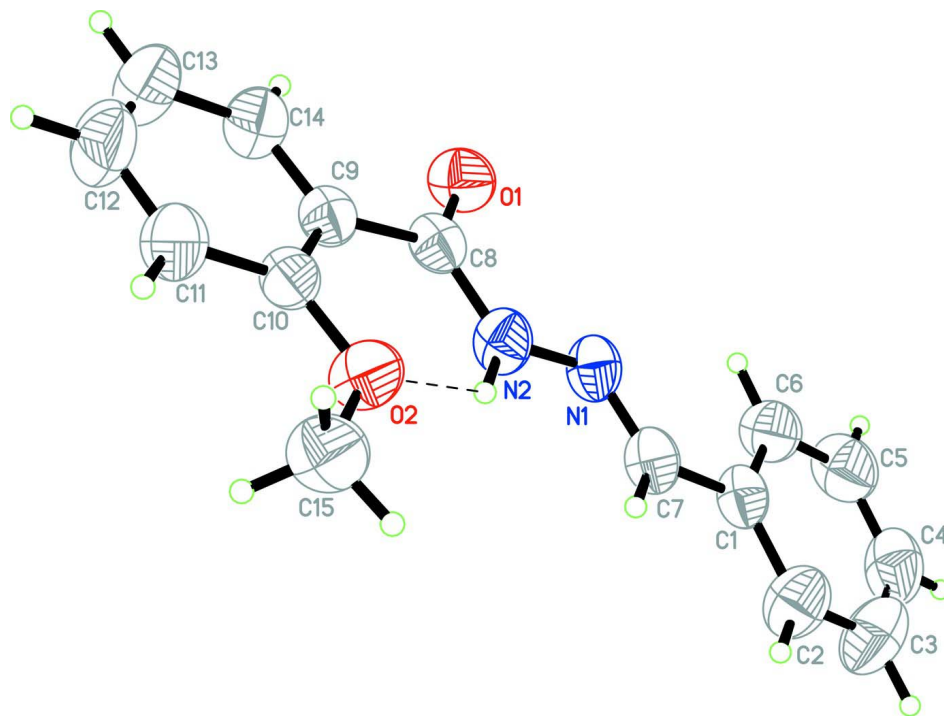
Applications of benzoylhydrazones are reported in medicinal and analytical chemistry (Tarafder *et al.*, 2002). Benzoylhydrazones having heterocyclic rings have been reported to have antiglycation (Taha *et al.*, 2014), anticonvulsant (Kucukguzel *et al.*, 2004), antiproliferative (Kucukguzel *et al.*, 2003), antifungal and anti-HIV activities (Pandeya *et al.*, 1999). Several benzoylhydrazones have also shown interesting bioactivities including antiinflammatory (Todeschini *et al.*, 1998), antibacterial (Kaymakcioglu *et al.*, 2006), antimalarial (Melnyk *et al.*, 2006) and anticancer (Terzioglu & Gursoy, 2003). Recently they have been suggested as an alternative in UV-laser desorption ionization (LDI) matrices for peptides analysis (Musharraf *et al.*, 2012). The structure of the title compound (Fig. 1) is similar to (*E*)-2-methoxy-*N'*-(2,4,6-trihydroxybenzylidene) benzohydrazide (Taha *et al.*, 2013), the difference residing in that the trihydroxyphenyl ring has been replaced by a non-substituted phenyl ring (C1–C6). The bond lengths and angle were found to be similar to the structurally related benzohydrazide derivatives reported in Taha *et al.*, 2013. The *E* configuration around the azomethine double bond is stabilized by a N2—H2A···O2 intramolecular interaction. The crystal structure is in turn stabilized by the intermolecular C7—H7A···O1 interaction to form chains along the *b* axis (Table 2 and Fig. 2).

S2. Experimental

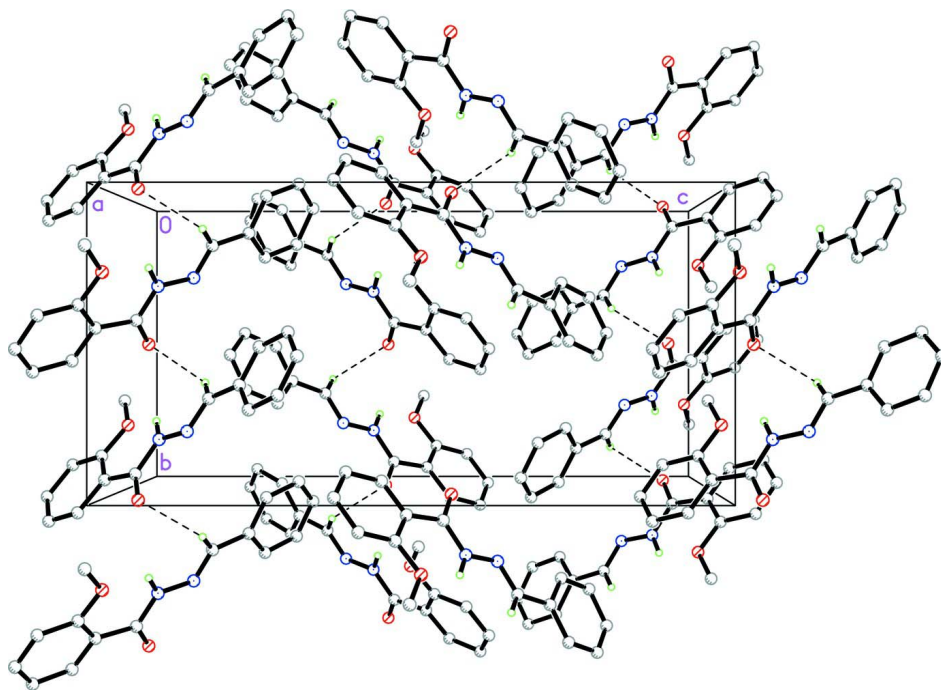
The title compound was synthesized by refluxing a mixture of 2 mmol of 2-methoxybenzohydrazide (0.332 g), 2 mmol benzaldehyde (0.212 g) and catalytic amount of acetic acid in methanol (20 ml) for 3 h. The progress of the reaction was monitored by TLC. After completion of the reaction, the solvent was evaporated by vacuum to afford crude product which was further recrystallized in methanol to afford needle like pure product in 88% yield (0.447 g). All the chemicals were purchased from sigma Aldrich Germany.

S3. Refinement

H atoms were positioned geometrically with C—H = 0.93/0.96 Å, N—H = 0.86 Å respectively, and constrained to ride on their parent atoms with $U_{\text{iso}}(\text{H}) = 1.2/1.5 U_{\text{eq}}(\text{CH})$.

**Figure 1**

The molecular structure of (I) with displacement ellipsoids drawn at 50% probability level.

**Figure 2**

The crystal packing of the title compound I. Only hydrogen atoms involved in hydrogen bonding are shown.

(E)-N'-Benzylidene-2-methoxybenzohydrazide*Crystal data*C₁₅H₁₄N₂O₂ $M_r = 254.28$ Orthorhombic, *Pbca* $a = 13.3135$ (9) Å $b = 9.9581$ (6) Å $c = 20.0278$ (14) Å $V = 2655.2$ (3) Å³ $Z = 8$ $F(000) = 1072$ $D_x = 1.272$ Mg m⁻³

Melting point = 449–451 K

Mo *K*α radiation, $\lambda = 0.71073$ Å

Cell parameters from 9899 reflections

 $\theta = 3.0$ – 25.5° $\mu = 0.09$ mm⁻¹ $T = 296$ K

Block, colourless

 $0.50 \times 0.48 \times 0.31$ mm*Data collection*Bruker SMART APEX CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 83.66 pixels mm⁻¹ ω scanAbsorption correction: multi-scan
(*SADABS*; Bruker, 2000) $T_{\min} = 0.94$, $T_{\max} = 0.97$

38941 measured reflections

2462 independent reflections

1819 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.049$ $\theta_{\max} = 25.5^\circ$, $\theta_{\min} = 3.0^\circ$ $h = -16 \rightarrow 16$ $k = -11 \rightarrow 12$ $l = -24 \rightarrow 24$ *Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.038$ $wR(F^2) = 0.103$ $S = 1.08$

2462 reflections

174 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.0361P)^2 + 0.849P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.13$ e Å⁻³ $\Delta\rho_{\min} = -0.14$ 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 (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.67310 (8)	-0.00143 (12)	0.55943 (6)	0.0569 (3)
O2	0.92308 (8)	0.22579 (11)	0.51387 (6)	0.0575 (3)
N1	0.69422 (9)	0.21778 (14)	0.63568 (6)	0.0481 (4)
N2	0.75847 (10)	0.18784 (14)	0.58377 (6)	0.0486 (3)

H2A	0.8086	0.2394	0.5750	0.058*
C1	0.65053 (12)	0.36836 (17)	0.72316 (7)	0.0466 (4)
C2	0.66885 (14)	0.4903 (2)	0.75366 (10)	0.0661 (5)
H2B	0.7240	0.5414	0.7408	0.079*
C3	0.60545 (17)	0.5369 (2)	0.80336 (11)	0.0797 (6)
H3A	0.6183	0.6190	0.8238	0.096*
C4	0.52403 (16)	0.4624 (2)	0.82254 (10)	0.0733 (6)
H4A	0.4806	0.4946	0.8552	0.088*
C5	0.50683 (15)	0.3407 (2)	0.79347 (10)	0.0701 (6)
H5A	0.4523	0.2891	0.8071	0.084*
C6	0.56950 (13)	0.29365 (18)	0.74411 (9)	0.0585 (5)
H6A	0.5570	0.2105	0.7247	0.070*
C9	0.80953 (11)	0.05112 (16)	0.48857 (7)	0.0434 (4)
C14	0.78281 (13)	-0.05488 (19)	0.44740 (9)	0.0587 (5)
H14A	0.7254	-0.1042	0.4573	0.070*
C13	0.83879 (16)	-0.0892 (2)	0.39229 (10)	0.0727 (6)
H13A	0.8196	-0.1609	0.3654	0.087*
C12	0.92326 (16)	-0.0164 (2)	0.37740 (10)	0.0730 (6)
H12A	0.9613	-0.0389	0.3401	0.088*
C11	0.95232 (14)	0.08876 (19)	0.41671 (9)	0.0605 (5)
H11A	1.0097	0.1374	0.4058	0.073*
C10	0.89683 (11)	0.12330 (16)	0.47268 (8)	0.0452 (4)
C15	1.00920 (14)	0.30448 (19)	0.49805 (10)	0.0662 (5)
H15A	1.0165	0.3749	0.5304	0.099*
H15B	1.0012	0.3431	0.4545	0.099*
H15C	1.0679	0.2485	0.4986	0.099*
C7	0.71491 (11)	0.32331 (17)	0.66852 (7)	0.0472 (4)
H7A	0.7719	0.3726	0.6576	0.057*
C8	0.74169 (11)	0.07632 (16)	0.54683 (7)	0.0431 (4)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0488 (7)	0.0567 (7)	0.0652 (8)	-0.0047 (6)	0.0131 (6)	0.0040 (6)
O2	0.0509 (7)	0.0553 (7)	0.0664 (8)	-0.0085 (5)	0.0210 (6)	-0.0075 (6)
N1	0.0414 (7)	0.0615 (9)	0.0413 (7)	0.0032 (6)	0.0088 (6)	0.0011 (7)
N2	0.0402 (7)	0.0594 (9)	0.0461 (7)	-0.0032 (6)	0.0126 (6)	-0.0014 (7)
C1	0.0436 (9)	0.0579 (10)	0.0383 (8)	0.0066 (8)	-0.0003 (7)	0.0039 (7)
C2	0.0573 (11)	0.0763 (13)	0.0649 (11)	-0.0044 (10)	0.0020 (9)	-0.0128 (10)
C3	0.0799 (14)	0.0884 (15)	0.0708 (13)	0.0113 (12)	0.0001 (12)	-0.0298 (12)
C4	0.0756 (14)	0.0940 (16)	0.0504 (11)	0.0258 (12)	0.0128 (10)	-0.0019 (11)
C5	0.0690 (13)	0.0747 (14)	0.0665 (12)	0.0094 (10)	0.0263 (10)	0.0095 (11)
C6	0.0602 (11)	0.0581 (10)	0.0574 (10)	0.0037 (8)	0.0167 (9)	0.0015 (9)
C9	0.0390 (8)	0.0476 (9)	0.0435 (9)	0.0065 (7)	0.0018 (7)	0.0047 (7)
C14	0.0561 (11)	0.0653 (11)	0.0547 (10)	-0.0044 (9)	0.0022 (8)	-0.0046 (9)
C13	0.0794 (14)	0.0804 (14)	0.0584 (12)	-0.0032 (12)	0.0064 (10)	-0.0198 (10)
C12	0.0764 (13)	0.0871 (15)	0.0556 (11)	0.0023 (12)	0.0211 (10)	-0.0119 (11)
C11	0.0577 (11)	0.0671 (12)	0.0569 (11)	0.0002 (9)	0.0193 (9)	0.0003 (9)

C10	0.0444 (9)	0.0465 (9)	0.0446 (9)	0.0079 (7)	0.0063 (7)	0.0031 (7)
C15	0.0567 (11)	0.0658 (12)	0.0762 (12)	-0.0137 (9)	0.0166 (9)	-0.0003 (10)
C7	0.0382 (8)	0.0613 (11)	0.0420 (8)	0.0012 (8)	0.0029 (7)	0.0044 (8)
C8	0.0357 (8)	0.0489 (9)	0.0447 (8)	0.0063 (7)	0.0019 (7)	0.0078 (7)

Geometric parameters (Å, °)

O1—C8	1.2236 (18)	C5—H5A	0.9300
O2—C10	1.3579 (19)	C6—H6A	0.9300
O2—C15	1.424 (2)	C9—C14	1.386 (2)
N1—C7	1.270 (2)	C9—C10	1.403 (2)
N1—N2	1.3790 (17)	C9—C8	1.497 (2)
N2—C8	1.353 (2)	C14—C13	1.375 (2)
N2—H2A	0.8600	C14—H14A	0.9300
C1—C6	1.376 (2)	C13—C12	1.371 (3)
C1—C2	1.381 (2)	C13—H13A	0.9300
C1—C7	1.461 (2)	C12—C11	1.366 (3)
C2—C3	1.385 (3)	C12—H12A	0.9300
C2—H2B	0.9300	C11—C10	1.386 (2)
C3—C4	1.368 (3)	C11—H11A	0.9300
C3—H3A	0.9300	C15—H15A	0.9600
C4—C5	1.364 (3)	C15—H15B	0.9600
C4—H4A	0.9300	C15—H15C	0.9600
C5—C6	1.376 (2)	C7—H7A	0.9300
C10—O2—C15	119.05 (13)	C13—C14—H14A	119.1
C7—N1—N2	115.77 (13)	C9—C14—H14A	119.1
C8—N2—N1	119.16 (13)	C12—C13—C14	119.22 (19)
C8—N2—H2A	120.4	C12—C13—H13A	120.4
N1—N2—H2A	120.4	C14—C13—H13A	120.4
C6—C1—C2	118.61 (16)	C11—C12—C13	120.82 (18)
C6—C1—C7	121.51 (16)	C11—C12—H12A	119.6
C2—C1—C7	119.86 (16)	C13—C12—H12A	119.6
C1—C2—C3	120.31 (19)	C12—C11—C10	120.36 (17)
C1—C2—H2B	119.8	C12—C11—H11A	119.8
C3—C2—H2B	119.8	C10—C11—H11A	119.8
C4—C3—C2	120.2 (2)	O2—C10—C11	122.73 (15)
C4—C3—H3A	119.9	O2—C10—C9	117.41 (13)
C2—C3—H3A	119.9	C11—C10—C9	119.86 (16)
C5—C4—C3	119.64 (18)	O2—C15—H15A	109.5
C5—C4—H4A	120.2	O2—C15—H15B	109.5
C3—C4—H4A	120.2	H15A—C15—H15B	109.5
C4—C5—C6	120.5 (2)	O2—C15—H15C	109.5
C4—C5—H5A	119.8	H15A—C15—H15C	109.5
C6—C5—H5A	119.8	H15B—C15—H15C	109.5
C5—C6—C1	120.70 (18)	N1—C7—C1	120.99 (15)
C5—C6—H6A	119.7	N1—C7—H7A	119.5
C1—C6—H6A	119.7	C1—C7—H7A	119.5

C14—C9—C10	117.90 (15)	O1—C8—N2	122.00 (14)
C14—C9—C8	115.89 (14)	O1—C8—C9	120.33 (15)
C10—C9—C8	126.21 (14)	N2—C8—C9	117.66 (14)
C13—C14—C9	121.84 (17)		
C7—N1—N2—C8	179.48 (14)	C12—C11—C10—O2	179.16 (17)
C6—C1—C2—C3	-1.2 (3)	C12—C11—C10—C9	-0.9 (3)
C7—C1—C2—C3	176.93 (17)	C14—C9—C10—O2	-179.12 (14)
C1—C2—C3—C4	-0.1 (3)	C8—C9—C10—O2	0.2 (2)
C2—C3—C4—C5	1.4 (3)	C14—C9—C10—C11	0.9 (2)
C3—C4—C5—C6	-1.4 (3)	C8—C9—C10—C11	-179.76 (15)
C4—C5—C6—C1	0.0 (3)	N2—N1—C7—C1	177.81 (13)
C2—C1—C6—C5	1.3 (3)	C6—C1—C7—N1	4.7 (2)
C7—C1—C6—C5	-176.83 (16)	C2—C1—C7—N1	-173.47 (16)
C10—C9—C14—C13	-0.4 (3)	N1—N2—C8—O1	-2.5 (2)
C8—C9—C14—C13	-179.76 (17)	N1—N2—C8—C9	176.67 (13)
C9—C14—C13—C12	-0.2 (3)	C14—C9—C8—O1	6.6 (2)
C14—C13—C12—C11	0.3 (3)	C10—C9—C8—O1	-172.75 (15)
C13—C12—C11—C10	0.3 (3)	C14—C9—C8—N2	-172.60 (14)
C15—O2—C10—C11	2.2 (2)	C10—C9—C8—N2	8.1 (2)
C15—O2—C10—C9	-177.76 (15)		

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

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N2—H2 <i>A</i> \cdots O2	0.86	1.96	2.6278 (17)	134
C7—H7 <i>A</i> \cdots O1 ⁱ	0.93	2.44	3.1690 (19)	135
C14—H14 <i>A</i> \cdots O1	0.93	2.39	2.730 (2)	101

Symmetry code: (i) $-x+3/2, y+1/2, z$.