

Acta Crystallographica Section E

## Structure Reports

Online

ISSN 1600-5368

**(E)-N'-(2-Hydroxy-4-methoxybenzylidene)isonicotinohydrazone monohydrate**San-Jun Peng<sup>a\*</sup> and Hai-Yun Hou<sup>b</sup>

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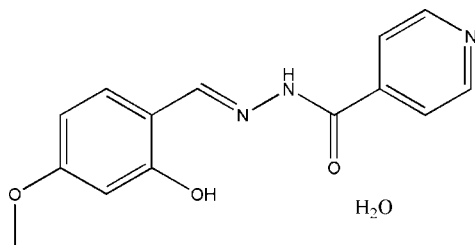
Received 12 September 2008; accepted 16 September 2008

Key indicators: single-crystal X-ray study;  $T = 298$  K; mean  $\sigma(\text{C}-\text{C}) = 0.002$  Å; R factor = 0.046;  $wR$  factor = 0.113; data-to-parameter ratio = 15.1.

The title compound,  $\text{C}_{14}\text{H}_{13}\text{N}_3\text{O}_3 \cdot \text{H}_2\text{O}$ , was prepared by the reaction of 4-methoxysalicylaldehyde and isonicotinohydrazone in ethanol. The Schiff base molecule is not planar and has an *E* configuration with respect to the methyldene unit. The dihedral angle between the benzene and pyridine rings is  $36.8(2)^\circ$ . In the molecule there is an intramolecular  $\text{O}-\text{H} \cdots \text{N}$  hydrogen bond involving the hydroxyl substituent and the N atom of the 2-hydroxy-4-methoxybenzylidene unit. In the crystal, the molecules are linked through intermolecular  $\text{O}-\text{H} \cdots \text{O}$ ,  $\text{O}-\text{H} \cdots \text{N}$  and  $\text{N}-\text{H} \cdots \text{O}$  hydrogen bonds, forming layers parallel to the *bc* plane.

## Related literature

For bond-length data, see: Allen *et al.* (1987). For background on the biological properties of hydrazones, see: El-Tabl *et al.* (2008); Chen *et al.* (2008); Alvarez *et al.* (2008); Ventura & Martins (2008); Kalinowski *et al.* (2008). For related structures, see: Peng & Hou (2008); Shan *et al.* (2008); Fun *et al.* (2008); Yehye *et al.* (2008); Ejsmont *et al.* (2008); Han *et al.* (2006); Lu *et al.* (2008).



## Experimental

## Crystal data

$\text{C}_{14}\text{H}_{13}\text{N}_3\text{O}_3 \cdot \text{H}_2\text{O}$   
 $M_r = 289.29$   
 Monoclinic,  $P2_1/c$   
 $a = 7.299(4)$  Å  
 $b = 12.537(6)$  Å  
 $c = 14.808(7)$  Å  
 $\beta = 96.281(8)^\circ$

$V = 1346.9(11)$  Å<sup>3</sup>  
 $Z = 4$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.11$  mm<sup>-1</sup>  
 $T = 298(2)$  K  
 $0.23 \times 0.23 \times 0.22$  mm

## Data collection

Bruker SMART 1000 CCD area-detector diffractometer  
 Absorption correction: multi-scan (SADABS; Bruker, 2001)  
 $T_{\min} = 0.976$ ,  $T_{\max} = 0.977$

7804 measured reflections  
 3041 independent reflections  
 2129 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.030$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.045$   
 $wR(F^2) = 0.112$   
 $S = 1.03$   
 3041 reflections  
 201 parameters  
 4 restraints

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\text{max}} = 0.18$  e Å<sup>-3</sup>  
 $\Delta\rho_{\text{min}} = -0.25$  e Å<sup>-3</sup>

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>
O1-H1...N1	0.82	1.92	2.644 (2)	146
O4-H4B...O2 <sup>i</sup>	0.853 (9)	2.072 (10)	2.924 (2)	176 (2)
O4-H4A...N3 <sup>ii</sup>	0.861 (9)	1.971 (10)	2.832 (2)	178 (2)
N2-H2...O4 <sup>iii</sup>	0.903 (9)	2.024 (11)	2.915 (2)	169 (2)

Symmetry codes: (i)  $-x + 1, -y + 1, -z + 1$ ; (ii)  $x, y + 1, z$ ; (iii)  $x, -y + \frac{1}{2}, z + \frac{1}{2}$ .

Data collection: SMART (Bruker, 2007); cell refinement: SAINT (Bruker, 2007); data reduction: SAINT; 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.

The corresponding author gratefully acknowledges Changsha University of Science and Technology for research grants.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SU2062).

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## supporting information

*Acta Cryst.* (2008). E64, o1996–o1997 [doi:10.1107/S1600536808029619]

**(E)-N'-(2-Hydroxy-4-methoxybenzylidene)isonicotinohydrazide monohydrate****San-Jun Peng and Hai-Yun Hou****S1. Comment**

Hydrazones derived from the reactions of aldehydes with hydrazides show potential biological properties (El-Tabl *et al.*, 2008; Chen *et al.*, 2008; Alvarez *et al.*, 2008; Ventura & Martins, 2008; Kalinowski *et al.*, 2008). In the last few years, a large number of hydrazones have been reported (Peng & Hou, 2008; Shan *et al.*, 2008; Fun *et al.*, 2008; Yehye *et al.*, 2008; Ejsmont *et al.*, 2008). As a continuous study, the crystal structure of the title compound, (I), is reported in this paper.

The molecular structure of compound (I) is illustrated in Fig. 1. It consists of a Schiff base molecule and a water molecule of crystallization. The C7=N1 bond length of 1.276 (2) Å indicates a typical C=N double bond. The Schiff base molecule has an E configuration with respect to the methyldene unit (C7=N1), as observed in similar compounds (Han *et al.*, 2006; Lu *et al.*, 2008). In the molecule there is an intramolecular O-H...N hydrogen bond involving the hydroxyl substituent and the N-atom of the 2-hydroxy-4-methoxybenzylidene moiety (Table 1). The dihedral angle between the benzene and pyridine rings is 36.8 (2)°, indicating the molecule is not planar. The bond lengths are in normal ranges (Allen *et al.*, 1987).

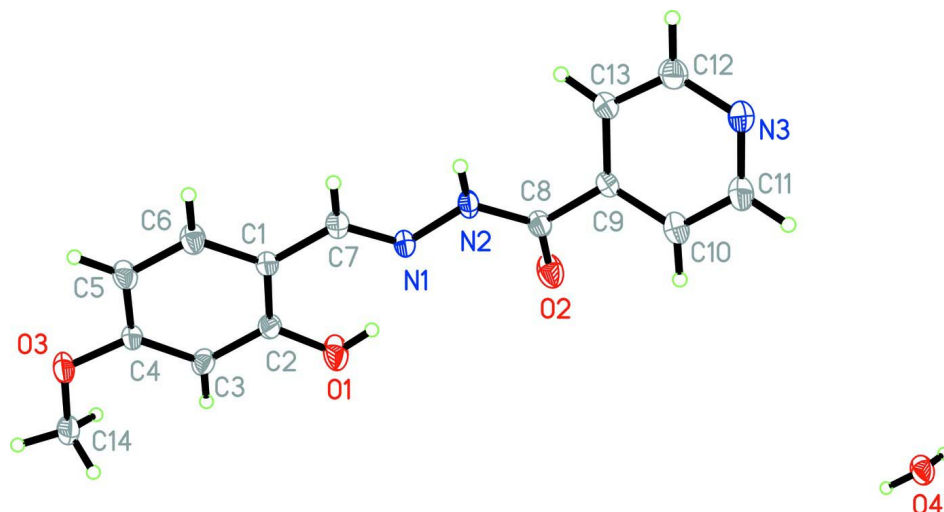
In the crystal structure, symmetry related molecules are linked through intermolecular O—H...O, O—H...N and N—H...O hydrogen bonds (Table 1), forming layers parallel to the *bc* plane (Fig. 2).

**S2. Experimental**

4-Methoxysalicylaldehyde (0.152 g, 1 mmol) was dissolved in 95% ethanol (50 ml), then isonicotinohydrazide (0.137 g, 1 mmol) was added slowly to the solution, and the mixture was heated at reflux with continuous stirring for 1 h. The solution was cooled to room temperature, yielding colorless crystallites. Recrystallization from a 95% ethanol yielded block-like single crystals of compound (I).

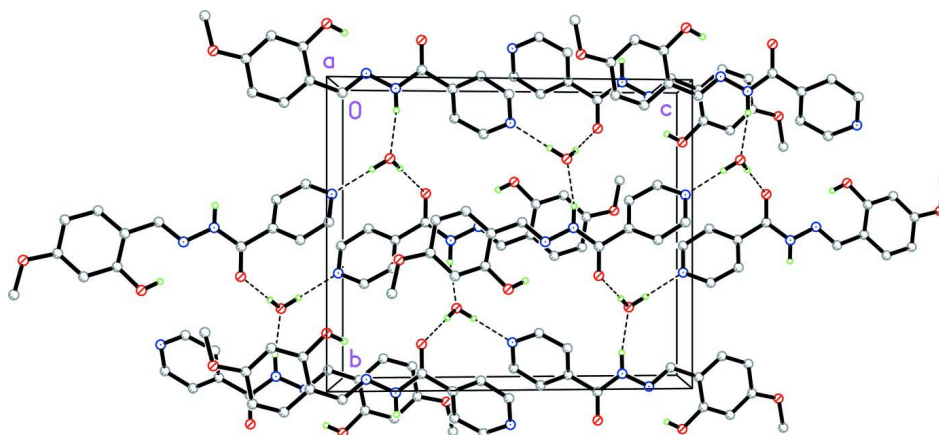
**S3. Refinement**

H-atoms H2, H4A and H4B were located in a difference Fourier map and refined isotropically, with N—H, O—H and H...H distances restrained to 0.90 (1), 0.85 (1) and 1.37 (2) Å, respectively, and with  $U_{\text{iso}}(\text{H})$  set at 0.08 Å<sup>2</sup>. The other H atoms were placed in calculated positions and treated as riding atoms with C—H = 0.93 - 0.96 Å, O—H = 0.82 Å, and  $i > U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$  and  $1.5U_{\text{eq}}(\text{O1 and C14})$ .



**Figure 1**

The molecular structure of compound (I), with 30% probability displacement ellipsoids for non-H atoms.



**Figure 2**

The crystal packing diagram of compound (I), viewed along the *a* axis. Hydrogen bonds are shown as dashed lines.

**(E)-N'-(2-Hydroxy-4-methoxybenzylidene)isonicotinohydrazide monohydrate**

*Crystal data*

$C_{14}H_{13}N_3O_3 \cdot H_2O$

$M_r = 289.29$

Monoclinic,  $P2_1/c$

Hall symbol:  $-P 2_1 ybc$

$a = 7.299$  (4) Å

$b = 12.537$  (6) Å

$c = 14.808$  (7) Å

$\beta = 96.281$  (8)°

$V = 1346.9$  (11) Å<sup>3</sup>

$Z = 4$

$F(000) = 608$

$D_x = 1.427$  Mg m<sup>-3</sup>

Mo  $K\alpha$  radiation,  $\lambda = 0.71073$  Å

Cell parameters from 1237 reflections

$\theta = 2.4$ – $24.5$ °

$\mu = 0.11$  mm<sup>-1</sup>

$T = 298$  K

Block, colorless

$0.23 \times 0.23 \times 0.22$  mm

*Data collection*

Bruker SMART 1000 CCD area-detector  
diffractometer  
Radiation source: fine-focus sealed tube  
Graphite monochromator  
 $\omega$  scans  
Absorption correction: multi-scan  
(*SADABS*; Bruker, 2001)  
 $T_{\min} = 0.976$ ,  $T_{\max} = 0.977$

7804 measured reflections  
3041 independent reflections  
2129 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.030$   
 $\theta_{\max} = 27.5^\circ$ ,  $\theta_{\min} = 2.1^\circ$   
 $h = -9 \rightarrow 9$   
 $k = -16 \rightarrow 15$   
 $l = -14 \rightarrow 19$

*Refinement*

Refinement on  $F^2$   
Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.046$   
 $wR(F^2) = 0.112$   
 $S = 1.03$   
3041 reflections  
201 parameters  
4 restraints  
Primary atom site location: structure-invariant  
direct methods

Secondary atom site location: difference Fourier  
map  
Hydrogen site location: inferred from  
neighbouring sites  
H atoms treated by a mixture of independent  
and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0459P)^2 + 0.2867P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} < 0.001$   
 $\Delta\rho_{\max} = 0.18 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.25 \text{ e } \text{\AA}^{-3}$

*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 ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.1869 (2)	0.17985 (9)	1.00835 (8)	0.0519 (4)
H1	0.2127	0.1524	0.9611	0.078*
O2	0.3338 (2)	0.13791 (9)	0.75220 (8)	0.0490 (4)
O3	0.07206 (17)	0.08252 (9)	1.30718 (7)	0.0407 (3)
O4	0.3910 (2)	0.75061 (10)	0.34205 (9)	0.0532 (4)
N1	0.29393 (19)	0.02521 (11)	0.90464 (9)	0.0348 (3)
N2	0.3324 (2)	-0.02124 (11)	0.82441 (9)	0.0339 (3)
N3	0.3613 (2)	-0.13153 (13)	0.50292 (10)	0.0496 (4)
C1	0.2257 (2)	-0.00223 (12)	1.05635 (10)	0.0307 (4)
C2	0.1816 (2)	0.10410 (13)	1.07253 (10)	0.0324 (4)
C3	0.1288 (2)	0.13524 (13)	1.15565 (10)	0.0342 (4)
H3	0.0991	0.2061	1.1657	0.041*
C4	0.1205 (2)	0.06050 (13)	1.22317 (10)	0.0312 (4)
C5	0.1644 (2)	-0.04517 (13)	1.20894 (11)	0.0370 (4)
H5	0.1592	-0.0953	1.2549	0.044*

C6	0.2154 (2)	-0.07474 (13)	1.12659 (11)	0.0382 (4)
H6	0.2442	-0.1458	1.1172	0.046*
C7	0.2761 (2)	-0.03892 (14)	0.97002 (11)	0.0355 (4)
H7	0.2962	-0.1114	0.9618	0.043*
C8	0.3435 (2)	0.04041 (13)	0.75139 (10)	0.0335 (4)
C9	0.3600 (2)	-0.01964 (12)	0.66533 (10)	0.0317 (4)
C10	0.2673 (3)	0.01868 (15)	0.58540 (11)	0.0417 (4)
H10	0.2037	0.0830	0.5847	0.050*
C11	0.2709 (3)	-0.03991 (16)	0.50692 (12)	0.0501 (5)
H11	0.2064	-0.0139	0.4538	0.060*
C12	0.4556 (3)	-0.16551 (15)	0.57962 (12)	0.0442 (5)
H12	0.5238	-0.2280	0.5776	0.053*
C13	0.4581 (2)	-0.11375 (13)	0.66161 (11)	0.0358 (4)
H13	0.5244	-0.1414	0.7136	0.043*
C14	0.0200 (3)	0.18921 (14)	1.32548 (12)	0.0460 (5)
H14A	0.1199	0.2366	1.3168	0.069*
H14B	-0.0087	0.1943	1.3871	0.069*
H14C	-0.0864	0.2086	1.2849	0.069*
H2	0.342 (3)	-0.0930 (8)	0.8225 (16)	0.080*
H4A	0.380 (3)	0.7860 (16)	0.3910 (10)	0.080*
H4B	0.476 (2)	0.7815 (17)	0.3165 (13)	0.080*

*Atomic displacement parameters (Å<sup>2</sup>)*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0907 (11)	0.0363 (7)	0.0306 (7)	-0.0014 (7)	0.0150 (7)	0.0072 (5)
O2	0.0739 (10)	0.0350 (7)	0.0413 (7)	0.0045 (6)	0.0206 (6)	0.0002 (5)
O3	0.0562 (8)	0.0416 (7)	0.0263 (6)	0.0017 (6)	0.0139 (5)	-0.0029 (5)
O4	0.0814 (11)	0.0405 (8)	0.0406 (8)	-0.0106 (7)	0.0193 (7)	-0.0080 (6)
N1	0.0376 (8)	0.0437 (8)	0.0241 (7)	-0.0035 (6)	0.0080 (6)	-0.0041 (6)
N2	0.0423 (8)	0.0373 (8)	0.0234 (7)	-0.0013 (6)	0.0093 (6)	-0.0038 (6)
N3	0.0564 (11)	0.0604 (11)	0.0340 (9)	-0.0128 (8)	0.0148 (7)	-0.0113 (7)
C1	0.0321 (9)	0.0351 (9)	0.0252 (8)	-0.0028 (7)	0.0045 (6)	-0.0011 (7)
C2	0.0401 (10)	0.0319 (8)	0.0248 (8)	-0.0056 (7)	0.0022 (7)	0.0034 (6)
C3	0.0454 (10)	0.0276 (8)	0.0297 (9)	-0.0012 (7)	0.0050 (7)	-0.0028 (7)
C4	0.0326 (9)	0.0382 (9)	0.0231 (8)	-0.0031 (7)	0.0048 (6)	-0.0021 (7)
C5	0.0497 (11)	0.0346 (9)	0.0279 (9)	0.0011 (8)	0.0095 (7)	0.0067 (7)
C6	0.0505 (11)	0.0308 (9)	0.0345 (9)	0.0045 (8)	0.0105 (8)	0.0007 (7)
C7	0.0401 (10)	0.0380 (9)	0.0289 (9)	-0.0008 (7)	0.0066 (7)	-0.0029 (7)
C8	0.0366 (9)	0.0356 (9)	0.0293 (9)	0.0004 (7)	0.0087 (7)	0.0002 (7)
C9	0.0349 (9)	0.0354 (9)	0.0262 (8)	-0.0042 (7)	0.0094 (6)	0.0010 (7)
C10	0.0488 (11)	0.0458 (10)	0.0313 (9)	0.0046 (8)	0.0081 (8)	0.0052 (8)
C11	0.0562 (13)	0.0653 (13)	0.0286 (10)	-0.0055 (10)	0.0042 (8)	0.0029 (9)
C12	0.0471 (11)	0.0433 (10)	0.0447 (11)	-0.0026 (8)	0.0166 (9)	-0.0090 (8)
C13	0.0364 (10)	0.0408 (9)	0.0312 (9)	-0.0027 (7)	0.0082 (7)	0.0002 (7)
C14	0.0585 (12)	0.0435 (10)	0.0387 (10)	-0.0069 (9)	0.0171 (8)	-0.0122 (8)

*Geometric parameters (Å, °)*

O1—C2	1.3471 (19)	C3—H3	0.9300
O1—H1	0.8200	C4—C5	1.385 (2)
O2—C8	1.224 (2)	C5—C6	1.365 (2)
O3—C4	1.3583 (19)	C5—H5	0.9300
O3—C14	1.425 (2)	C6—H6	0.9300
O4—H4A	0.861 (9)	C7—H7	0.9300
O4—H4B	0.853 (9)	C8—C9	1.496 (2)
N1—C7	1.276 (2)	C9—C10	1.383 (2)
N1—N2	1.3791 (19)	C9—C13	1.384 (2)
N2—C8	1.339 (2)	C10—C11	1.377 (2)
N2—H2	0.903 (9)	C10—H10	0.9300
N3—C11	1.329 (3)	C11—H11	0.9300
N3—C12	1.332 (2)	C12—C13	1.375 (2)
C1—C6	1.390 (2)	C12—H12	0.9300
C1—C2	1.398 (2)	C13—H13	0.9300
C1—C7	1.444 (2)	C14—H14A	0.9600
C2—C3	1.386 (2)	C14—H14B	0.9600
C3—C4	1.376 (2)	C14—H14C	0.9600
C2—O1—H1	109.5	N1—C7—H7	119.1
C4—O3—C14	117.81 (13)	C1—C7—H7	119.1
H4A—O4—H4B	106.4 (17)	O2—C8—N2	124.03 (15)
C7—N1—N2	115.77 (14)	O2—C8—C9	121.38 (14)
C8—N2—N1	119.21 (14)	N2—C8—C9	114.52 (14)
C8—N2—H2	122.6 (15)	C10—C9—C13	117.94 (15)
N1—N2—H2	118.1 (15)	C10—C9—C8	118.47 (15)
C11—N3—C12	116.82 (15)	C13—C9—C8	123.55 (15)
C6—C1—C2	117.59 (14)	C11—C10—C9	118.96 (17)
C6—C1—C7	119.62 (15)	C11—C10—H10	120.5
C2—C1—C7	122.77 (14)	C9—C10—H10	120.5
O1—C2—C3	117.41 (15)	N3—C11—C10	123.61 (17)
O1—C2—C1	121.76 (14)	N3—C11—H11	118.2
C3—C2—C1	120.83 (14)	C10—C11—H11	118.2
C4—C3—C2	119.47 (15)	N3—C12—C13	123.90 (18)
C4—C3—H3	120.3	N3—C12—H12	118.1
C2—C3—H3	120.3	C13—C12—H12	118.1
O3—C4—C3	124.30 (15)	C12—C13—C9	118.71 (16)
O3—C4—C5	114.93 (14)	C12—C13—H13	120.6
C3—C4—C5	120.77 (15)	C9—C13—H13	120.6
C6—C5—C4	119.09 (15)	O3—C14—H14A	109.5
C6—C5—H5	120.5	O3—C14—H14B	109.5
C4—C5—H5	120.5	H14A—C14—H14B	109.5
C5—C6—C1	122.25 (16)	O3—C14—H14C	109.5
C5—C6—H6	118.9	H14A—C14—H14C	109.5
C1—C6—H6	118.9	H14B—C14—H14C	109.5
N1—C7—C1	121.88 (15)		

*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>
O1—H1···N1	0.82	1.92	2.644 (2)	146
O4—H4B···O2 <sup>i</sup>	0.85 (1)	2.07 (1)	2.924 (2)	176 (2)
O4—H4A···N3 <sup>ii</sup>	0.86 (1)	1.97 (1)	2.832 (2)	178 (2)
N2—H2···O4 <sup>iii</sup>	0.90 (1)	2.02 (1)	2.915 (2)	169 (2)

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