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### 2-(6-Hydroxy-1-benzofuran-3-yl)acetamide

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Key indicators: single-crystal X-ray study; T = 293 K; mean  $\sigma$ (C–C) = 0.002 Å; R factor = 0.033; wR factor = 0.081; data-to-parameter ratio = 10.7.

In the title compound,  $C_{10}H_9NO_3$ , the dihedral angle between the benzofuran ring system (r.m.s. deviation for the non-H atoms = 0.009 Å) and the -C-C(O)-N- segment is 83.76 (1)°. In the crystal, molecules are linked by  $N-H\cdots O$ and  $O-H\cdots O$  hydrogen bonds, generating (001) sheets, which feature C(4) and C(10) chains.

#### **Related literature**

For a related structure and background to benzofurans, see: Arunakumar *et al.* (2014).



#### Experimental

Crystal data  $C_{10}H_9NO_3$  $M_r = 191.18$ 

Orthorhombic,  $P2_12_12_1$ a = 5.0939 (3) Å b = 9.3629 (5) Å c = 18.7422 (10) Å  $V = 893.88 (9) \text{ Å}^3$ Z = 4

#### Data collection

Bruker APEXII CCD diffractometer Absorption correction: multi-scan (SADABS; Bruker, 2009)  $T_{min} = 0.770, T_{max} = 0.808$ 

#### Refinement

$$\begin{split} R[F^2 > 2\sigma(F^2)] &= 0.033 \\ wR(F^2) &= 0.081 \\ S &= 1.14 \\ 1459 \text{ reflections} \\ 136 \text{ parameters} \\ H \text{ atoms treated by a mixture of} \\ \text{independent and constrained} \\ \text{refinement} \end{split}$$

# Cu $K\alpha$ radiation

organic compounds

 $\mu = 0.89 \text{ mm}^{-1}$  T = 293 K $0.36 \times 0.29 \times 0.24 \text{ mm}$ 

8714 measured reflections 1459 independent reflections 1446 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.036$ 

 $\begin{array}{l} \Delta \rho_{\rm max} = 0.15 \mbox{ e } {\rm \AA}^{-3} \\ \Delta \rho_{\rm min} = -0.28 \mbox{ e } {\rm \AA}^{-3} \\ \mbox{Absolute structure: Flack (1983),} \\ 1927 \mbox{ Friedel pairs} \\ \mbox{Absolute structure parameter:} \\ -0.2 \mbox{ (2)} \end{array}$ 

# Table 1Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$\begin{array}{c} O3 - HO3 \cdots O1^{i} \\ N1 - H1N \cdots O3^{ii} \\ N1 - H2N \cdots O1^{iii} \end{array}$	0.82	1.92	2.7006 (16)	158
	0.92 (3)	2.08 (3)	2.969 (2)	162 (2)
	0.92 (3)	2.03 (3)	2.8958 (18)	156 (2)

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

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT-Plus* (Bruker, 2009); data reduction: *SAINT-Plus* and *XPREP* (Bruker, 2009); 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: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB7172).

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# supplementary materials

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## 2-(6-Hydroxy-1-benzofuran-3-yl)acetamide

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#### 1. Comment

As part of our ongoing structural studies of functionalsied benzofuran ring systems at the C-2 position (Arunakumar et al., 2014), we now describe the structure of the title compound, (I).

In the title compound,  $C_{10}H_9NO_3$ , the benzofuran ring is almost planar ((r.m.s. deviation for the non-H atoms = 0.009 Å) (Fig. 1). The dihedral angle between the two planes defined by the benzofuran ring and the -C-C(O)-N segment is 96.24 (1)°. Compared to this, the dihedral angle between the benzofuran ring and the -C-C-N o segment in the related structure (Arunakumar *et al.*, 2014) is 2.85 (1)°. In the crystal structure, the molecules are linked to one another through strong N1-H1N···O3 and N1-H2N···O1 hydrogen bonds generating C(4) and C(10) chains (Fig. 2). The molecules are further connected into C(10) chains through strong O3-HO3···O1 hydrogen bonds forming helical structure (Fig. 3). Linking of molecules in zig zag pattern through N1-H1N···O3 H-bonds is shown in Fig. 4.

#### 2. Experimental

(6-Hydroxy-1-benzofuran-3-yl)acetic acid (0.015 mmol) was taken in a round bottom flask containing *N*,*N*-dimethyl formamide (15 ml)·To this 1, 1-carbonydiimadazole (0.023 mmol) was added at 273 K. The reaction mixture was stirred for 45 minutes. Ammonium acetate (0.046 mmol) and triethyl amine (1 ml) were added, the reaction mixture was stirred at room temperature overnight. The completion of the reaction was monitored by thin layer chromatography. After completion of the reaction mixture was diluted with ethylacetate and the organic layer was washed with water followed by brine solution. The organic layer was dried over anhydrous sodium sulfate and concentrated to give the crude product. It was further purified by column chromatography by using Ethyl acetate and petroleum ether (8:2) as eluent.

Colourless prisms were obtained from the solvent system: ethyl acetate / methanol (4:1) at room temperature.

#### 3. Refinement

The H atoms were positioned with idealized geometry using a riding model with C—H = 0.93–0.97 Å and O—H = 0.82 Å. The H atoms of the NH<sub>2</sub> group were located in a difference map and later refined freely. The isotropic displacement parameters for all H atoms were set to 1.2 times  $U_{eq}$  (Carbon) and 1.5 times  $U_{eq}$  (Oxygen)..

#### **Computing details**

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT-Plus* (Bruker, 2009); data reduction: *SAINT-Plus* and *XPREP* (Bruker, 2009); 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: *SHELXL97* (Sheldrick, 2008).





Molecular structure of the title compound, showing displacement ellipsoids drawn at the 50% probability level.



### Figure 2

Linking of molecules in the crystal structure through N—H···O hydrogen bonds into C(4) and C(10) chains. H-atoms not involved in H-bonding are omitted for clarity.



### Figure 3

Formation of helical structure through O—H…O hydrogen bonds.



#### Figure 4

Zig-zag pattern observed in the crystal structure.

#### 2-(6-Hydroxy-1-benzofuran-3-yl)acetamide

Crystal data

C<sub>10</sub>H<sub>9</sub>NO<sub>3</sub>  $M_r = 191.18$ Orthorhombic,  $P2_12_12_1$ Hall symbol: P 2ac 2ab a = 5.0939 (3) Å b = 9.3629 (5) Å c = 18.7422 (10) Å V = 893.88 (9) Å<sup>3</sup> Z = 4F(000) = 400

#### Data collection

Bruker APEXII CCD diffractometer Radiation source: fine-focus sealed tube Graphite monochromator  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Bruker, 2009)  $T_{\min} = 0.770, T_{\max} = 0.808$  Prism  $D_x = 1.421 \text{ Mg m}^{-3}$ Melting point: 533 K Cu Ka radiation,  $\lambda = 1.54178 \text{ Å}$ Cell parameters from 1234 reflections  $\theta = 4.7-64.3^{\circ}$   $\mu = 0.89 \text{ mm}^{-1}$  T = 293 KPrism, colourless  $0.36 \times 0.29 \times 0.24 \text{ mm}$ 

8714 measured reflections 1459 independent reflections 1446 reflections with  $I > 2\sigma(I)$  $R_{int} = 0.036$  $\theta_{max} = 64.3^\circ, \theta_{min} = 4.7^\circ$  $h = -5 \rightarrow 5$  $k = -10 \rightarrow 10$  $l = -21 \rightarrow 21$  Refinement

Refinement on $F^2$ Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.033$ wR(F^2) = 0.081	H atoms treated by a mixture of independent and constrained refinement
S = 1.14 1459 reflections	$w = 1/[\sigma^2(F_o^2) + (0.0488P)^2 + 0.1043P]$ where $P = (F_o^2 + 2F_c^2)/3$
136 parameters	$(\Delta/\sigma)_{\rm max} < 0.001$
0 restraints	$\Delta \rho_{\rm max} = 0.15 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	$\Delta \rho_{\min} = -0.28 \text{ e} \text{ Å}^{-3}$ Absolute structure: Flack (1983), 1927 Friedel
Secondary atom site location: difference Fourier map	pairs Absolute structure parameter: -0.2 (2)

#### Special details

**Geometry**. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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  $F^2 > 2\sigma(F^2)$  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	$U_{ m iso}$ */ $U_{ m eq}$	
H2N	0.766 (5)	0.410 (3)	0.6264 (14)	0.072 (7)*	
H1N	0.587 (5)	0.324 (3)	0.5742 (14)	0.069 (7)*	
C1	0.4600 (3)	1.02627 (16)	0.55047 (8)	0.0361 (4)	
C2	0.6218 (3)	0.90706 (18)	0.54019 (9)	0.0404 (4)	
H2	0.7494	0.9092	0.5047	0.049*	
C3	0.5949 (3)	0.78677 (18)	0.58188 (8)	0.0402 (4)	
H3	0.7026	0.7079	0.5747	0.048*	
C4	0.4029 (3)	0.78533 (16)	0.63518 (7)	0.0334 (3)	
C5	0.2448 (3)	0.90548 (17)	0.64295 (8)	0.0369 (4)	
C6	0.2657 (3)	1.02782 (17)	0.60204 (9)	0.0398 (4)	
H6	0.1562	1.1061	0.6088	0.048*	
C7	0.1208 (4)	0.75102 (19)	0.72418 (8)	0.0452 (4)	
H7	0.0293	0.7101	0.7619	0.054*	
C8	0.3173 (3)	0.68572 (17)	0.68946 (8)	0.0370 (4)	
C9	0.4300 (3)	0.54149 (18)	0.70437 (8)	0.0406 (4)	
H9A	0.3482	0.5023	0.7468	0.049*	
H9B	0.6166	0.5506	0.7136	0.049*	
C10	0.3882 (3)	0.44027 (15)	0.64264 (8)	0.0331 (3)	
N1	0.5982 (3)	0.38614 (19)	0.61193 (9)	0.0492 (4)	
01	0.1637 (2)	0.41077 (13)	0.62188 (6)	0.0433 (3)	
O2	0.0692 (2)	0.88620 (13)	0.69762 (6)	0.0473 (3)	
O3	0.4914 (2)	1.14862 (11)	0.51112 (6)	0.0454 (3)	
HO3	0.5821	1.1314	0.4760	0.068*	

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
C1	0.0386 (8)	0.0371 (8)	0.0326 (7)	-0.0049 (6)	-0.0004 (6)	-0.0042 (6)
C2	0.0363 (9)	0.0486 (9)	0.0364 (7)	-0.0001 (7)	0.0078 (7)	-0.0003 (6)
C3	0.0371 (9)	0.0440 (8)	0.0396 (8)	0.0048 (7)	0.0066 (7)	-0.0023 (6)
C4	0.0306 (8)	0.0402 (8)	0.0294 (7)	-0.0038 (6)	0.0006 (6)	-0.0059 (6)
C5	0.0334 (8)	0.0444 (8)	0.0328 (7)	-0.0052 (7)	0.0050 (6)	-0.0093 (6)
C6	0.0381 (9)	0.0395 (8)	0.0417 (8)	0.0015 (7)	0.0042 (7)	-0.0081 (7)
C7	0.0499 (10)	0.0505 (9)	0.0353 (8)	-0.0090 (8)	0.0103 (7)	-0.0037 (7)
C8	0.0375 (9)	0.0449 (8)	0.0287 (7)	-0.0085 (7)	0.0000 (7)	-0.0056 (6)
C9	0.0411 (9)	0.0495 (9)	0.0312 (7)	-0.0058 (7)	-0.0054 (7)	0.0026 (6)
C10	0.0296 (8)	0.0377 (7)	0.0320 (7)	-0.0027 (6)	-0.0021 (6)	0.0070 (6)
N1	0.0320 (8)	0.0597 (9)	0.0559 (9)	-0.0004(7)	0.0012 (7)	-0.0097 (8)
O1	0.0290 (6)	0.0567 (7)	0.0444 (6)	-0.0025 (5)	-0.0029 (5)	-0.0082 (5)
O2	0.0476 (7)	0.0508 (7)	0.0434 (6)	-0.0008 (6)	0.0187 (5)	-0.0079 (5)
O3	0.0537 (8)	0.0405 (6)	0.0419 (6)	-0.0003 (5)	0.0081 (5)	0.0008 (5)

Atomic displacement parameters  $(Å^2)$ 

Geometric parameters (Å, °)

C1-03	1.3717 (19)	C7—C8	1.341 (2)
C1—C6	1.383 (2)	С7—О2	1.385 (2)
C1—C2	1.401 (2)	С7—Н7	0.9300
С2—С3	1.378 (2)	C8—C9	1.494 (2)
С2—Н2	0.9300	C9—C10	1.511 (2)
C3—C4	1.398 (2)	С9—Н9А	0.9700
С3—Н3	0.9300	С9—Н9В	0.9700
C4—C5	1.391 (2)	C10—O1	1.2390 (19)
C4—C8	1.447 (2)	C10—N1	1.317 (2)
С5—О2	1.372 (2)	N1—H2N	0.92 (3)
С5—С6	1.382 (2)	N1—H1N	0.92 (3)
С6—Н6	0.9300	О3—НО3	0.8200
O3—C1—C6	116.76 (14)	С8—С7—Н7	123.7
O3—C1—C2	121.52 (14)	O2—C7—H7	123.7
C6—C1—C2	121.69 (15)	C7—C8—C4	105.80 (15)
C3—C2—C1	120.98 (14)	C7—C8—C9	127.46 (16)
С3—С2—Н2	119.5	C4—C8—C9	126.73 (15)
С1—С2—Н2	119.5	C8—C9—C10	111.69 (12)
C2—C3—C4	118.87 (15)	С8—С9—Н9А	109.3
С2—С3—Н3	120.6	С10—С9—Н9А	109.3
С4—С3—Н3	120.6	C8—C9—H9B	109.3
C5—C4—C3	118.16 (14)	С10—С9—Н9В	109.3
C5—C4—C8	105.84 (13)	H9A—C9—H9B	107.9
C3—C4—C8	136.00 (15)	O1—C10—N1	121.80 (15)
O2—C5—C6	124.99 (15)	O1—C10—C9	120.70 (14)
O2—C5—C4	110.44 (14)	N1—C10—C9	117.50 (15)
C6—C5—C4	124.56 (14)	C10—N1—H2N	121.9 (16)
C5—C6—C1	115.73 (15)	C10—N1—H1N	122.0 (17)
С5—С6—Н6	122.1	H2N—N1—H1N	116 (2)

C1—C6—H6	122.1	C5—O2—C7	105.37 (13)
C8—C7—O2	112.56 (14)	C1—O3—HO3	109.5
$\begin{array}{c} 03 - C1 - C2 - C3 \\ C6 - C1 - C2 - C3 \\ C1 - C2 - C3 - C4 \\ C2 - C3 - C4 - C5 \\ C2 - C3 - C4 - C5 \\ C3 - C4 - C5 - 02 \\ C3 - C4 - C5 - 02 \\ C3 - C4 - C5 - C6 \\ C8 - C4 - C5 - C6 \\ 02 - C5 - C6 - C1 \end{array}$	177.50 (15)	O2-C7-C8-C4	-0.22 (19)
	-0.7 (3)	O2-C7-C8-C9	178.51 (14)
	-0.2 (2)	C5-C4-C8-C7	0.25 (17)
	0.9 (2)	C3-C4-C8-C7	179.59 (18)
	-178.39 (17)	C5-C4-C8-C9	-178.49 (14)
	-179.68 (13)	C3-C4-C8-C9	0.8 (3)
	-0.21 (17)	C7-C8-C9-C10	116.36 (19)
	-0.8 (2)	C4-C8-C9-C10	-65.2 (2)
	178.69 (15)	C8-C9-C10-O1	-59.9 (2)
	178.69 (15)	C8-C9-C10-N1	119.67 (16)
C4C5C6C1	-0.1 (2)	C6—C5—O2—C7	-178.81 (16)
O3C1C6C5	-177.48 (14)	C4—C5—O2—C7	0.08 (17)
C2C1C6C5	0.8 (2)	C8—C7—O2—C5	0.09 (18)

### Hydrogen-bond geometry (Å, °)

D—H···A	D—H	Н…А	D····A	<i>D</i> —H··· <i>A</i>
03—H03…01 <sup>i</sup>	0.82	1.92	2.7006 (16)	158
N1—H1 <i>N</i> ···O3 <sup>ii</sup>	0.92 (3)	2.08 (3)	2.969 (2)	162 (2)
N1—H2N····O1 <sup>iii</sup>	0.92 (3)	2.03 (3)	2.8958 (18)	156 (2)

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