

## [3-(Propenamido)phenyl]boronic acid

Dan Zhang,<sup>a</sup> Laura E. Harrington,<sup>b\*</sup> Hiroo Tanaka<sup>a</sup> and Robert Pelton<sup>a</sup>

<sup>a</sup>Department of Chemical Engineering, JHE-136, McMaster University, Hamilton, Ontario, Canada L8S 4L7, and <sup>b</sup>Department of Chemistry, McMaster University, 1280 Main Street W., Hamilton, Ontario, Canada L8S 4M1

Correspondence e-mail: harrin@mcmaster.ca

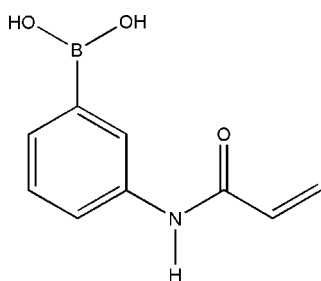
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Key indicators: single-crystal X-ray study;  $T = 295$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.038;  $wR$  factor = 0.108; data-to-parameter ratio = 9.6.

The title compound,  $\text{C}_9\text{H}_{10}\text{BNO}_3$ , was synthesized for investigation as a potential precursor for use in affinity chromatography and glucose sensing. Weak  $\text{N}-\text{H}\cdots\text{O}$  and  $\text{O}-\text{H}\cdots\text{O}$  hydrogen-bonding interactions are observed in the molecular structure. The dihedral angle between the benzene ring and the  $\text{BO}_2$  plane is  $35.1(2)^\circ$ .

### Related literature

For related literature, see: Zhang *et al.* (2007); Chen *et al.* (2006); Deutsch & Osoling (1949).



### Experimental

#### Crystal data

$\text{C}_9\text{H}_{10}\text{BNO}_3$   
 $M_r = 190.99$   
 Orthorhombic,  $Fdd2$   
 $a = 18.626(4)$  Å  
 $b = 42.112(8)$  Å  
 $c = 5.0536(10)$  Å

$V = 3963.9(13)$  Å<sup>3</sup>  
 $Z = 16$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.09$  mm<sup>-1</sup>  
 $T = 295(2)$  K  
 $0.43 \times 0.18 \times 0.12$  mm

#### Data collection

Bruker APEXII CCD diffractometer  
 Absorption correction: numerical (*APEX2*; Bruker, 2006)  
 $T_{\min} = 0.87$ ,  $T_{\max} = 0.99$

14806 measured reflections  
 1657 independent reflections  
 1279 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.035$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.039$   
 $wR(F^2) = 0.108$   
 $S = 1.04$   
 1657 reflections  
 172 parameters  
 1 restraint

H atoms treated by a mixture of independent and constrained refinement

$\Delta\rho_{\max} = 0.17$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.12$  e Å<sup>-3</sup>

**Table 1**

Hydrogen-bond geometry (Å, °).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{O2}-\text{H2A}\cdots\text{O2}^{\text{i}}$	0.77 (7)	2.02 (7)	2.759 (3)	163 (7)
$\text{O2}-\text{H2B}\cdots\text{O3}^{\text{ii}}$	0.77 (8)	2.10 (9)	2.748 (3)	143 (7)
$\text{O3}-\text{H3A}\cdots\text{O2}^{\text{iii}}$	0.75 (8)	2.24 (8)	2.748 (3)	126 (7)
$\text{O3}-\text{H3B}\cdots\text{O3}^{\text{i}}$	0.87 (5)	1.88 (5)	2.746 (3)	173 (6)
$\text{N1}-\text{H1A}\cdots\text{O1}^{\text{iv}}$	0.93 (3)	1.98 (3)	2.911 (3)	172 (2)

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

Data collection: *APEX2* (Bruker, 2006); cell refinement: *APEX2*; data reduction: *APEX2*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 2000) and *Mercury* (Macrae *et al.*, 2006); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: KJ2069).

### References

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**supplementary materials**

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### [3-(Propenamido)phenyl]boronic acid

D. Zhang, L. E. Harrington, H. Tanaka and R. Pelton

#### Comment

The title compound is a monomer containing a boronate group, which can form complexes with *cis* diols at pH values above 9 (Deutsch & Osoling, 1949). This interaction can be employed for affinity chromatography and glucose sensing (Chen *et al.*, 2006).

The molecular structure of the title compound is illustrated in Fig.1. Weak N—H···O and O—H···O hydrogen bonding interactions are observed in the molecular structure which help to stabilize the crystal structure and forming an infinite three-dimensional network, as illustrated in the packing diagram displayed in Fig. 2. The dihedral angle between the benzene ring and the plane composed of O2—B1—O3 is 35.12(0.23)°.

#### Experimental

The synthesis of (I) was described previously (Zhang *et al.*, 2007). Single crystals suitable for X-ray analysis were obtained by slow evaporation of a water solution at room temperature.

#### Refinement

Friedel pairs were merged prior to refinement using *SHELXTL* (Bruker, 2000). The positions of the H atoms were determined using the difference map and were refined isotropically. The H atoms on O2 and O3 were disordered over 2 sites with 50% occupancy in each site. The disordered H atoms were refined with one common  $U_{iso}$ .

#### Figures

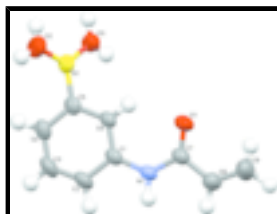


Fig. 1. The molecular structure of (I), showing the atom-labelling scheme. Displacement ellipsoids are drawn at the 50% probability level. H atoms are represented by circles of arbitrary size. Only one component of the disorder is shown.

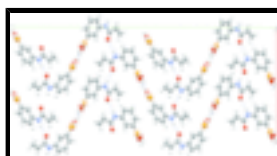


Fig. 2. The crystal packing of (II), viewed along the *c* axis. Only one component of the disorder is shown. H-bonds are shown as dashed lines.

## [3-(Propenamido)phenyl]boronic acid

### Crystal data

C<sub>9</sub>H<sub>10</sub>BNO<sub>3</sub>

*M<sub>r</sub>* = 190.99

Orthorhombic, *Fdd2*

Hall symbol: F 2 -2d

*a* = 18.626 (4) Å

*b* = 42.112 (8) Å

*c* = 5.0536 (10) Å

*V* = 3963.9 (13) Å<sup>3</sup>

*Z* = 16

*F*<sub>000</sub> = 1600

*D<sub>x</sub>* = 1.280 Mg m<sup>-3</sup>

Mo *K*α radiation

λ = 0.71073 Å

Cell parameters from 2853 reflections

θ = 2.4–25.1°

μ = 0.09 mm<sup>-1</sup>

*T* = 295 (2) K

Needle, colourless

0.43 × 0.18 × 0.12 mm

### Data collection

Bruker APEXII CCD  
diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

*T* = 295(2) K

φ and ω scans

Absorption correction: numerical  
(face correction with APEX2; Bruker, 2006)

*T<sub>min</sub>* = 0.87, *T<sub>max</sub>* = 0.99

14806 measured reflections

1657 independent reflections

1279 reflections with *I* > 2σ(*I*)

*R<sub>int</sub>* = 0.035

θ<sub>max</sub> = 30.5°

θ<sub>min</sub> = 1.9°

*h* = -26→25

*k* = -59→59

*l* = -7→2

### Refinement

Refinement on *F*<sup>2</sup>

Least-squares matrix: full

*R* [*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.039

*wR* (*F*<sup>2</sup>) = 0.108

*S* = 1.04

1657 reflections

172 parameters

1 restraint

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

$$w = 1/[\sigma^2(F_o^2) + (0.0606P)^2 + 0.5557P]$$

where *P* = (*F<sub>o</sub>*<sup>2</sup> + 2*F<sub>c</sub>*<sup>2</sup>)/3

(Δ/σ)<sub>max</sub> = 0.007

Δρ<sub>max</sub> = 0.17 e Å<sup>-3</sup>

Δρ<sub>min</sub> = -0.12 e Å<sup>-3</sup>

Extinction correction: none

*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  $F^2 > 2\sigma(F^2)$  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>	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
O1	0.34232 (7)	0.11569 (4)	1.1454 (4)	0.0653 (5)	
O2	0.43076 (12)	0.01158 (6)	0.3738 (4)	0.0779 (6)	
H2A	0.466 (3)	0.0024 (17)	0.394 (16)	0.082 (10)*	0.50
H2B	0.413 (4)	0.0131 (15)	0.237 (17)	0.082 (10)*	0.50
O3	0.43531 (9)	0.01566 (4)	0.8312 (3)	0.0569 (4)	
H3A	0.414 (4)	0.0220 (15)	0.944 (17)	0.082 (10)*	0.50
H3B	0.478 (3)	0.0072 (14)	0.827 (14)	0.082 (10)*	0.50
N1	0.23312 (9)	0.10547 (4)	0.9628 (3)	0.0476 (4)	
H1A	0.1859 (16)	0.1131 (6)	0.949 (6)	0.072 (8)*	
B1	0.39944 (13)	0.02160 (6)	0.6027 (5)	0.0469 (5)	
C1	0.24915 (10)	0.08023 (4)	0.7901 (4)	0.0442 (4)	
C2	0.19502 (12)	0.07111 (5)	0.6107 (5)	0.0567 (5)	
H2	0.1493 (13)	0.0830 (5)	0.624 (6)	0.059 (6)*	
C3	0.20685 (14)	0.04743 (6)	0.4318 (5)	0.0633 (6)	
H3	0.1704 (15)	0.0419 (6)	0.324 (7)	0.068 (7)*	
C4	0.27236 (13)	0.03202 (5)	0.4223 (4)	0.0543 (5)	
H4	0.2783 (13)	0.0162 (6)	0.288 (6)	0.060 (7)*	
C5	0.32638 (10)	0.03966 (4)	0.6037 (4)	0.0447 (4)	
C6	0.31372 (10)	0.06372 (5)	0.7880 (4)	0.0433 (4)	
H6	0.3479 (12)	0.0696 (5)	0.914 (5)	0.055 (6)*	
C7	0.27804 (10)	0.12142 (5)	1.1262 (4)	0.0478 (4)	
C8	0.24262 (14)	0.14649 (6)	1.2856 (6)	0.0646 (6)	
H8	0.1923 (18)	0.1488 (7)	1.258 (8)	0.085 (9)*	
C9	0.2770 (2)	0.16236 (8)	1.4653 (8)	0.0938 (11)	
H9A	0.2548 (17)	0.1772 (8)	1.613 (9)	0.101 (10)*	
H9B	0.322 (2)	0.1570 (8)	1.511 (9)	0.105 (12)*	

*Atomic displacement parameters ( $\text{\AA}^2$ )*

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
O1	0.0357 (7)	0.0729 (9)	0.0873 (12)	0.0017 (6)	-0.0151 (8)	-0.0233 (9)
O2	0.0824 (15)	0.1155 (17)	0.0359 (9)	0.0226 (11)	0.0051 (9)	-0.0134 (10)

## supplementary materials

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O3	0.0593 (10)	0.0778 (11)	0.0334 (7)	0.0152 (8)	-0.0019 (6)	-0.0005 (7)
N1	0.0334 (8)	0.0568 (9)	0.0526 (9)	0.0036 (7)	-0.0099 (7)	-0.0050 (8)
B1	0.0551 (12)	0.0549 (11)	0.0307 (8)	0.0020 (9)	0.0008 (8)	-0.0002 (9)
C1	0.0394 (9)	0.0505 (10)	0.0427 (9)	-0.0035 (8)	-0.0090 (7)	0.0036 (8)
C2	0.0463 (11)	0.0635 (12)	0.0602 (13)	-0.0005 (10)	-0.0215 (10)	-0.0013 (11)
C3	0.0607 (13)	0.0686 (14)	0.0606 (13)	-0.0068 (11)	-0.0281 (11)	-0.0065 (11)
C4	0.0648 (13)	0.0559 (11)	0.0421 (10)	-0.0038 (10)	-0.0128 (9)	-0.0034 (9)
C5	0.0507 (10)	0.0511 (10)	0.0324 (8)	-0.0022 (8)	-0.0017 (7)	0.0052 (8)
C6	0.0380 (9)	0.0550 (11)	0.0370 (9)	-0.0014 (8)	-0.0066 (7)	-0.0001 (7)
C7	0.0381 (9)	0.0530 (10)	0.0522 (10)	-0.0016 (8)	-0.0076 (8)	-0.0007 (9)
C8	0.0525 (13)	0.0653 (13)	0.0760 (17)	0.0114 (11)	-0.0125 (12)	-0.0159 (12)
C9	0.081 (2)	0.097 (2)	0.104 (3)	0.0237 (17)	-0.0317 (18)	-0.049 (2)

### Geometric parameters ( $\text{\AA}$ , $^\circ$ )

O1—C7	1.225 (2)	C2—C3	1.364 (4)
O2—B1	1.363 (3)	C2—H2	0.99 (2)
O2—H2A	0.76 (7)	C3—C4	1.383 (4)
O2—H2B	0.77 (8)	C3—H3	0.90 (3)
O3—B1	1.357 (3)	C4—C5	1.399 (3)
O3—H3A	0.74 (8)	C4—H4	0.96 (3)
O3—H3B	0.87 (5)	C5—C6	1.396 (3)
N1—C7	1.354 (3)	C6—H6	0.93 (3)
N1—C1	1.407 (3)	C7—C8	1.483 (3)
N1—H1A	0.94 (3)	C8—C9	1.297 (4)
B1—C5	1.559 (3)	C8—H8	0.95 (3)
C1—C6	1.389 (3)	C9—H9A	1.06 (4)
C1—C2	1.409 (3)	C9—H9B	0.89 (4)
B1—O2—H2A	114 (6)	C4—C3—H3	121.6 (17)
B1—O2—H2B	123 (6)	C3—C4—C5	120.3 (2)
H2A—O2—H2B	122 (8)	C3—C4—H4	116.9 (15)
B1—O3—H3A	109 (6)	C5—C4—H4	122.9 (15)
B1—O3—H3B	120 (5)	C6—C5—C4	118.86 (18)
H3A—O3—H3B	131 (7)	C6—C5—B1	120.24 (17)
C7—N1—C1	128.46 (17)	C4—C5—B1	120.90 (19)
C7—N1—H1A	117.1 (18)	C1—C6—C5	120.95 (16)
C1—N1—H1A	114.2 (18)	C1—C6—H6	117.0 (14)
O3—B1—O2	117.0 (2)	C5—C6—H6	122.0 (14)
O3—B1—C5	121.1 (2)	O1—C7—N1	123.7 (2)
O2—B1—C5	121.8 (2)	O1—C7—C8	122.2 (2)
C6—C1—N1	124.49 (16)	N1—C7—C8	114.19 (18)
C6—C1—C2	118.57 (19)	C9—C8—C7	121.8 (3)
N1—C1—C2	116.93 (18)	C9—C8—H8	122 (2)
C3—C2—C1	120.7 (2)	C7—C8—H8	116 (2)
C3—C2—H2	123.6 (16)	C8—C9—H9A	127.2 (19)
C1—C2—H2	115.7 (15)	C8—C9—H9B	121 (2)
C2—C3—C4	120.6 (2)	H9A—C9—H9B	109 (3)
C2—C3—H3	117.8 (18)		
C7—N1—C1—C6	10.7 (3)	O3—B1—C5—C4	-145.3 (2)

C7—N1—C1—C2	-170.2 (2)	O2—B1—C5—C4	35.9 (3)
C6—C1—C2—C3	-2.6 (3)	N1—C1—C6—C5	-177.70 (18)
N1—C1—C2—C3	178.2 (2)	C2—C1—C6—C5	3.2 (3)
C1—C2—C3—C4	-0.3 (4)	C4—C5—C6—C1	-0.9 (3)
C2—C3—C4—C5	2.6 (4)	B1—C5—C6—C1	179.45 (19)
C3—C4—C5—C6	-2.0 (3)	C1—N1—C7—O1	0.3 (3)
C3—C4—C5—B1	177.6 (2)	C1—N1—C7—C8	-178.9 (2)
O3—B1—C5—C6	34.3 (3)	O1—C7—C8—C9	-4.2 (4)
O2—B1—C5—C6	-144.5 (2)	N1—C7—C8—C9	174.9 (3)

*Hydrogen-bond geometry (Å, °)*

<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>
O2—H2A $\cdots$ O2 <sup>i</sup>	0.77 (7)	2.02 (7)	2.759 (3)	163 (7)
O2—H2B $\cdots$ O3 <sup>ii</sup>	0.77 (8)	2.10 (9)	2.748 (3)	143 (7)
O3—H3A $\cdots$ O2 <sup>iii</sup>	0.75 (8)	2.24 (8)	2.748 (3)	126 (7)
O3—H3B $\cdots$ O3 <sup>i</sup>	0.87 (5)	1.88 (5)	2.746 (3)	173 (6)
N1—H1A $\cdots$ O1 <sup>iv</sup>	0.93 (3)	1.98 (3)	2.911 (3)	172 (2)

Symmetry codes: (i)  $-x+1, -y, z$ ; (ii)  $x, y, z-1$ ; (iii)  $x, y, z+1$ ; (iv)  $x-1/4, -y+1/4, z-1/4$ .

Fig. 1

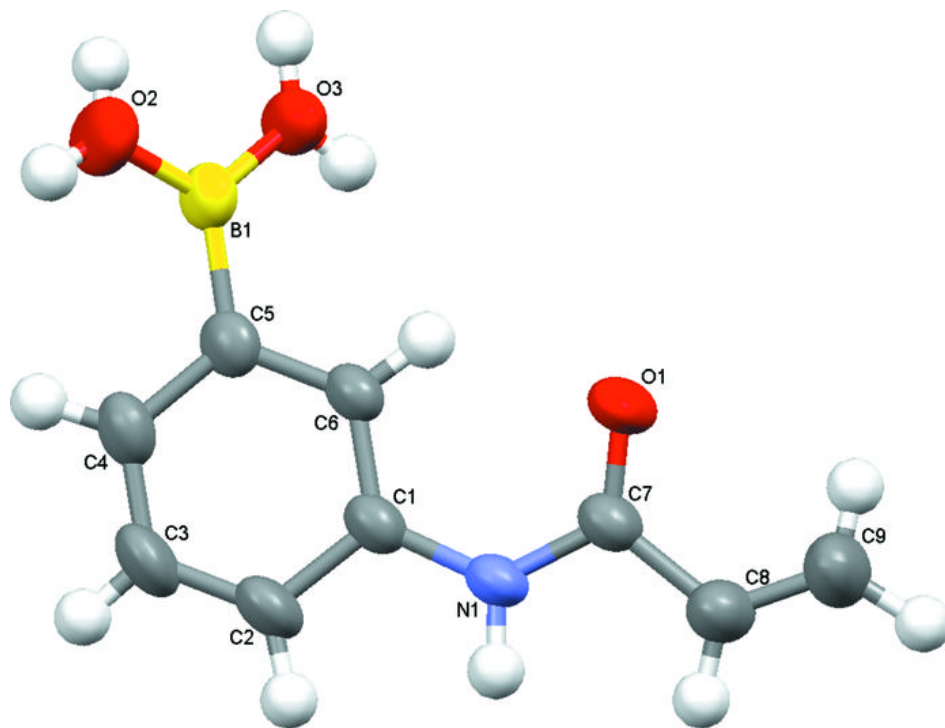




Fig. 2

