

Acta Crystallographica Section E

## Structure Reports

Online

ISSN 1600-5368

## 4-(3-Methyl-5-phenyl-1H-pyrazol-1-yl)-benzenesulfonamide

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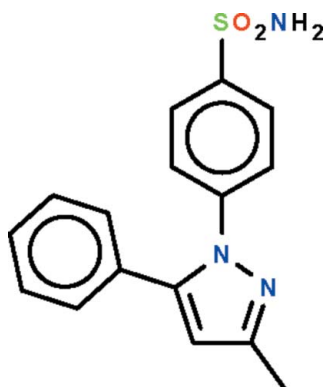
Received 11 August 2011; accepted 13 August 2011

Key indicators: single-crystal X-ray study;  $T = 100$  K; mean  $\sigma(\text{C}-\text{C}) = 0.003$  Å;  $R$  factor = 0.041;  $wR$  factor = 0.115; data-to-parameter ratio = 15.1.

With respect to the planar five-membered ring of the title compound,  $\text{C}_{16}\text{H}_{15}\text{N}_3\text{O}_2\text{S}$ , the phenyl ring is aligned at  $47.0(1)^\circ$  and the phenylene ring at  $37.6(1)^\circ$ . The amino group has the N atom in a pyramidal geometry; the group is a hydrogen-bond donor to the sulfonyl O atom of one molecule and to the pyrazole N atom of another molecule, resulting in the formation of a layer parallel to the  $bc$  plane.

## Related literature

For the synthesis, see: Gosselin *et al.* (2006); Organ & Mayer (2003).



## Experimental

## Crystal data

$\text{C}_{16}\text{H}_{15}\text{N}_3\text{O}_2\text{S}$   
 $M_r = 313.37$   
Monoclinic,  $C2/c$   
 $a = 28.2545(8)$  Å  
 $b = 11.9135(4)$  Å  
 $c = 9.3739(3)$  Å  
 $\beta = 91.016(3)^\circ$

$V = 3154.85(17)$  Å<sup>3</sup>  
 $Z = 8$   
Cu  $K\alpha$  radiation  
 $\mu = 1.91$  mm<sup>-1</sup>  
 $T = 100$  K  
 $0.30 \times 0.03 \times 0.03$  mm

## Data collection

Agilent SuperNova Dual  
diffractometer with an Atlas  
detector  
Absorption correction: multi-scan  
(*CrysAlis PRO*; Agilent, 2010)  
 $T_{\min} = 0.598$ ,  $T_{\max} = 0.945$

6579 measured reflections  
3137 independent reflections  
2689 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.035$

## Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$   
 $wR(F^2) = 0.115$   
 $S = 1.03$   
3137 reflections  
208 parameters

H atoms treated by a mixture of  
independent and constrained  
refinement  
 $\Delta\rho_{\max} = 0.33$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.51$  e Å<sup>-3</sup>

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{N3}^{\text{i}}$	0.92 (2)	1.98 (2)	2.878 (2)	164 (2)
$\text{N1}-\text{H2}\cdots\text{O1}^{\text{ii}}$	0.86 (2)	2.07 (2)	2.930 (2)	177 (2)

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

Data collection: *CrysAlis PRO* (Agilent, 2010); cell refinement: *CrysAlis PRO*; data reduction: *CrysAlis PRO*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *pubCIF* (Westrip, 2010).

We thank King Abdulaziz University and the University of Malaya for supporting this study.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5609).

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## supporting information

*Acta Cryst.* (2011). E67, o2426 [doi:10.1107/S1600536811032855]

**4-(3-Methyl-5-phenyl-1*H*-pyrazol-1-yl)benzenesulfonamide**

**Abdullah M. Asiri, Hassan M. Faidallah, Abdulrahman O. Al-Youbi, Salem A. Basaif and Seik Weng Ng**

**S1. Comment**

We are examining the medicinal properties of phenylpyrazolones of which the 4-benzenesulfamide derivative (Scheme I) is expected to show enhanced activity. As the inhibitory activity against cyclooxygenase-1 and cyclooxygenase-2 of the title compound (Scheme I) has been claimed in a number of patents, other researchers have attempted its synthesis in order to increase yield (Gosselin *et al.*, 2006; Organ & Mayer, 2003). With respect to the planar five-membered ring, the phenyl ring is aligned at 47.0 (1)° and the phenylene ring at 37.6 (1)°. The amino group is hydrogen bond donor to the sulfonyl O atom of one molecule and to the pyrazolyl N atom of another molecule to result in the formation of a layer parallel to the bc plane.

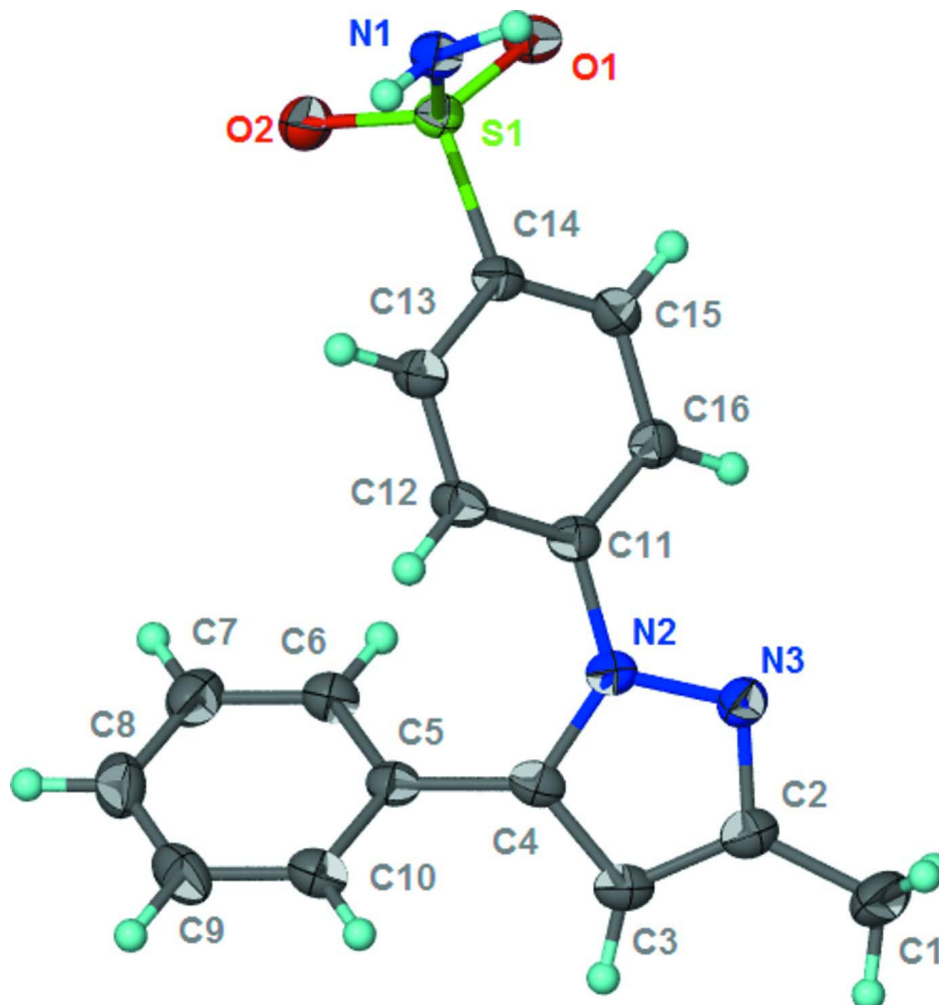
**S2. Experimental**

1-Phenylbutan-1,3-dione (10 mmol) and 4-hydrazinobenzenesulfonamide hydrochloride (10 mmol) were heated in ethanol (50 ml) for 4 h; water was added to precipitate the product, which was collected and recrystallized from ethanol as light yellow crystals; m.p. 471–472 K.

**S3. Refinement**

Carbon bound H-atoms were placed in calculated positions [C–H 0.95 to 0.98 Å,  $U_{\text{iso}}(\text{H})$  1.2–1.5 $U_{\text{eq}}(\text{C})$ ] and were included in the refinement in the riding model approximation.

The amino H-atoms were located in a difference Fourier map and were freely refined.

**Figure 1**

Anisotropic displacement ellipsoid plot (Barbour, 2001) of  $C_{16}H_{15}N_3O_2S$  at the 70% probability level; hydrogen atoms are drawn as spheres of arbitrary radius.

#### 4-(3-Methyl-5-phenyl-1H-pyrazol-1-yl)benzenesulfonamide

##### Crystal data

$C_{16}H_{15}N_3O_2S$

$M_r = 313.37$

Monoclinic,  $C2/c$

Hall symbol:  $-C 2yc$

$a = 28.2545 (8) \text{ \AA}$

$b = 11.9135 (4) \text{ \AA}$

$c = 9.3739 (3) \text{ \AA}$

$\beta = 91.016 (3)^\circ$

$V = 3154.85 (17) \text{ \AA}^3$

$Z = 8$

$F(000) = 1312$

$D_x = 1.320 \text{ Mg m}^{-3}$

Cu  $K\alpha$  radiation,  $\lambda = 1.54184 \text{ \AA}$

Cell parameters from 2773 reflections

$\theta = 3.1\text{--}74.2^\circ$

$\mu = 1.91 \text{ mm}^{-1}$

$T = 100 \text{ K}$

Prism, light-yellow

$0.30 \times 0.03 \times 0.03 \text{ mm}$

*Data collection*

Agilent SuperNova Dual  
 diffractometer with an Atlas detector  
 Radiation source: SuperNova (Cu) X-ray  
 Source  
 Mirror monochromator  
 Detector resolution: 10.4041 pixels mm<sup>-1</sup>  
 $\omega$  scans  
 Absorption correction: multi-scan  
 (CrysAlis PRO; Agilent, 2010)

$T_{\min} = 0.598$ ,  $T_{\max} = 0.945$   
 6579 measured reflections  
 3137 independent reflections  
 2689 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.035$   
 $\theta_{\max} = 74.4^\circ$ ,  $\theta_{\min} = 3.1^\circ$   
 $h = -35 \rightarrow 31$   
 $k = -12 \rightarrow 14$   
 $l = -11 \rightarrow 9$

*Refinement*

Refinement on  $F^2$   
 Least-squares matrix: full  
 $R[F^2 > 2\sigma(F^2)] = 0.041$   
 $wR(F^2) = 0.115$   
 $S = 1.03$   
 3137 reflections  
 208 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 atoms treated by a mixture of independent  
 and constrained refinement  
 $w = 1/[\sigma^2(F_o^2) + (0.0629P)^2 + 1.4812P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.001$   
 $\Delta\rho_{\max} = 0.33 \text{ e } \text{\AA}^{-3}$   
 $\Delta\rho_{\min} = -0.51 \text{ e } \text{\AA}^{-3}$

*Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters ( $\text{\AA}^2$ )*

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.690362 (14)	0.48353 (4)	0.65383 (4)	0.01752 (14)
O1	0.67646 (4)	0.41033 (11)	0.76803 (13)	0.0220 (3)
O2	0.69504 (5)	0.60135 (11)	0.68142 (14)	0.0252 (3)
N1	0.65210 (5)	0.46753 (14)	0.52785 (16)	0.0191 (3)
H1	0.6414 (8)	0.395 (2)	0.516 (2)	0.026 (6)*
H2	0.6583 (8)	0.5054 (18)	0.452 (2)	0.017 (5)*
N2	0.88441 (5)	0.35478 (13)	0.48169 (15)	0.0189 (3)
N3	0.89694 (5)	0.24415 (13)	0.47817 (17)	0.0217 (3)
C1	0.96762 (7)	0.13366 (18)	0.4272 (3)	0.0341 (5)
H1A	0.9686	0.0956	0.5199	0.051*
H1B	1.0000	0.1470	0.3954	0.051*
H1C	0.9508	0.0864	0.3573	0.051*
C2	0.94240 (6)	0.24326 (17)	0.4411 (2)	0.0234 (4)
C3	0.95933 (6)	0.35272 (17)	0.42301 (19)	0.0225 (4)
H3	0.9904	0.3741	0.3974	0.027*
C4	0.92187 (6)	0.42307 (16)	0.44972 (17)	0.0193 (4)
C5	0.91994 (6)	0.54696 (16)	0.44973 (18)	0.0192 (4)
C6	0.89996 (6)	0.60729 (16)	0.5616 (2)	0.0229 (4)
H6	0.8878	0.5683	0.6413	0.028*
C7	0.89779 (7)	0.72353 (17)	0.5573 (2)	0.0284 (4)
H7	0.8840	0.7639	0.6335	0.034*
C8	0.91585 (7)	0.78079 (17)	0.4409 (2)	0.0307 (5)
H8	0.9138	0.8603	0.4365	0.037*
C9	0.93685 (7)	0.72183 (18)	0.3313 (2)	0.0292 (4)

H9	0.9499	0.7612	0.2531	0.035*
C10	0.93889 (6)	0.60550 (17)	0.3355 (2)	0.0235 (4)
H10	0.9533	0.5656	0.2600	0.028*
C11	0.83692 (6)	0.38175 (15)	0.51674 (18)	0.0188 (4)
C12	0.81362 (6)	0.47000 (16)	0.44808 (19)	0.0214 (4)
H12	0.8285	0.5097	0.3733	0.026*
C13	0.76862 (6)	0.49917 (16)	0.48990 (19)	0.0213 (4)
H13	0.7529	0.5611	0.4465	0.026*
C14	0.74629 (6)	0.43750 (15)	0.59607 (18)	0.0183 (4)
C15	0.76849 (6)	0.34505 (15)	0.65805 (18)	0.0193 (4)
H15	0.7523	0.3004	0.7256	0.023*
C16	0.81447 (6)	0.31858 (15)	0.62039 (19)	0.0194 (4)
H16	0.8305	0.2576	0.6652	0.023*

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

	$U^{11}$	$U^{22}$	$U^{33}$	$U^{12}$	$U^{13}$	$U^{23}$
S1	0.0180 (2)	0.0196 (2)	0.0150 (2)	-0.00041 (15)	0.00220 (16)	-0.00277 (15)
O1	0.0219 (6)	0.0280 (7)	0.0161 (6)	0.0002 (5)	0.0038 (5)	-0.0002 (5)
O2	0.0254 (7)	0.0217 (7)	0.0287 (7)	-0.0010 (5)	0.0032 (5)	-0.0067 (5)
N1	0.0192 (7)	0.0221 (8)	0.0159 (7)	-0.0009 (6)	0.0013 (6)	0.0001 (6)
N2	0.0169 (7)	0.0212 (8)	0.0187 (7)	0.0003 (6)	0.0012 (6)	-0.0010 (6)
N3	0.0201 (7)	0.0195 (8)	0.0256 (8)	0.0009 (6)	0.0010 (6)	-0.0027 (6)
C1	0.0271 (10)	0.0280 (11)	0.0474 (13)	0.0041 (8)	0.0067 (9)	-0.0085 (9)
C2	0.0198 (9)	0.0270 (10)	0.0234 (9)	0.0012 (7)	0.0000 (7)	-0.0044 (7)
C3	0.0179 (8)	0.0283 (10)	0.0212 (9)	-0.0012 (7)	0.0017 (7)	-0.0037 (7)
C4	0.0180 (8)	0.0249 (9)	0.0149 (8)	-0.0034 (7)	0.0001 (6)	-0.0002 (7)
C5	0.0148 (8)	0.0236 (9)	0.0190 (9)	-0.0015 (7)	-0.0018 (6)	0.0007 (7)
C6	0.0207 (9)	0.0264 (10)	0.0218 (9)	-0.0014 (7)	0.0010 (7)	0.0004 (7)
C7	0.0248 (9)	0.0269 (10)	0.0335 (11)	0.0031 (8)	-0.0015 (8)	-0.0050 (8)
C8	0.0288 (10)	0.0219 (10)	0.0410 (12)	0.0007 (8)	-0.0067 (8)	0.0047 (9)
C9	0.0278 (10)	0.0302 (10)	0.0293 (10)	-0.0067 (8)	-0.0037 (8)	0.0098 (8)
C10	0.0206 (9)	0.0289 (10)	0.0211 (9)	-0.0038 (7)	0.0004 (7)	0.0009 (7)
C11	0.0161 (8)	0.0223 (9)	0.0179 (8)	-0.0009 (7)	-0.0001 (6)	-0.0023 (7)
C12	0.0189 (9)	0.0279 (10)	0.0175 (9)	-0.0022 (7)	0.0014 (7)	0.0054 (7)
C13	0.0186 (8)	0.0254 (9)	0.0199 (9)	-0.0003 (7)	-0.0003 (7)	0.0036 (7)
C14	0.0171 (8)	0.0210 (9)	0.0169 (8)	-0.0025 (7)	0.0005 (6)	-0.0031 (7)
C15	0.0220 (9)	0.0189 (9)	0.0171 (8)	-0.0028 (7)	0.0028 (7)	-0.0016 (7)
C16	0.0209 (9)	0.0192 (9)	0.0182 (8)	0.0003 (7)	0.0010 (6)	-0.0004 (7)

*Geometric parameters (Å, °)*

S1—O2	1.4330 (13)	C6—C7	1.387 (3)
S1—O1	1.4407 (13)	C6—H6	0.9500
S1—N1	1.5983 (15)	C7—C8	1.391 (3)
S1—C14	1.7666 (18)	C7—H7	0.9500
N1—H1	0.92 (2)	C8—C9	1.387 (3)
N1—H2	0.86 (2)	C8—H8	0.9500

N2—N3	1.365 (2)	C9—C10	1.388 (3)
N2—C4	1.372 (2)	C9—H9	0.9500
N2—C11	1.424 (2)	C10—H10	0.9500
N3—C2	1.337 (2)	C11—C16	1.391 (2)
C1—C2	1.494 (3)	C11—C12	1.392 (3)
C1—H1A	0.9800	C12—C13	1.382 (3)
C1—H1B	0.9800	C12—H12	0.9500
C1—H1C	0.9800	C13—C14	1.397 (2)
C2—C3	1.400 (3)	C13—H13	0.9500
C3—C4	1.376 (3)	C14—C15	1.390 (3)
C3—H3	0.9500	C15—C16	1.389 (2)
C4—C5	1.477 (3)	C15—H15	0.9500
C5—C10	1.393 (3)	C16—H16	0.9500
C5—C6	1.398 (3)		
O2—S1—O1	118.97 (8)	C7—C6—H6	119.7
O2—S1—N1	108.01 (8)	C5—C6—H6	119.7
O1—S1—N1	106.68 (8)	C6—C7—C8	119.71 (19)
O2—S1—C14	106.21 (8)	C6—C7—H7	120.1
O1—S1—C14	107.27 (8)	C8—C7—H7	120.1
N1—S1—C14	109.49 (8)	C9—C8—C7	120.03 (19)
S1—N1—H1	114.6 (14)	C9—C8—H8	120.0
S1—N1—H2	113.6 (14)	C7—C8—H8	120.0
H1—N1—H2	117.8 (19)	C8—C9—C10	120.20 (18)
N3—N2—C4	111.46 (14)	C8—C9—H9	119.9
N3—N2—C11	117.98 (14)	C10—C9—H9	119.9
C4—N2—C11	130.56 (16)	C9—C10—C5	120.36 (18)
C2—N3—N2	105.35 (15)	C9—C10—H10	119.8
C2—C1—H1A	109.5	C5—C10—H10	119.8
C2—C1—H1B	109.5	C16—C11—C12	120.91 (16)
H1A—C1—H1B	109.5	C16—C11—N2	118.85 (16)
C2—C1—H1C	109.5	C12—C11—N2	120.24 (16)
H1A—C1—H1C	109.5	C13—C12—C11	119.26 (16)
H1B—C1—H1C	109.5	C13—C12—H12	120.4
N3—C2—C3	110.85 (16)	C11—C12—H12	120.4
N3—C2—C1	119.44 (18)	C12—C13—C14	119.97 (17)
C3—C2—C1	129.70 (17)	C12—C13—H13	120.0
C4—C3—C2	106.24 (16)	C14—C13—H13	120.0
C4—C3—H3	126.9	C15—C14—C13	120.57 (16)
C2—C3—H3	126.9	C15—C14—S1	121.16 (13)
N2—C4—C3	106.09 (16)	C13—C14—S1	118.22 (14)
N2—C4—C5	124.32 (16)	C16—C15—C14	119.42 (16)
C3—C4—C5	129.57 (16)	C16—C15—H15	120.3
C10—C5—C6	119.01 (18)	C14—C15—H15	120.3
C10—C5—C4	119.05 (16)	C15—C16—C11	119.67 (16)
C6—C5—C4	121.94 (16)	C15—C16—H16	120.2
C7—C6—C5	120.64 (18)	C11—C16—H16	120.2

C4—N2—N3—C2	-1.13 (19)	C6—C5—C10—C9	-1.6 (3)
C11—N2—N3—C2	179.13 (15)	C4—C5—C10—C9	178.87 (16)
N2—N3—C2—C3	1.0 (2)	N3—N2—C11—C16	37.9 (2)
N2—N3—C2—C1	179.95 (17)	C4—N2—C11—C16	-141.76 (18)
N3—C2—C3—C4	-0.5 (2)	N3—N2—C11—C12	-142.14 (17)
C1—C2—C3—C4	-179.3 (2)	C4—N2—C11—C12	38.2 (3)
N3—N2—C4—C3	0.86 (19)	C16—C11—C12—C13	3.9 (3)
C11—N2—C4—C3	-179.44 (16)	N2—C11—C12—C13	-176.02 (16)
N3—N2—C4—C5	-177.79 (15)	C11—C12—C13—C14	-2.5 (3)
C11—N2—C4—C5	1.9 (3)	C12—C13—C14—C15	-1.5 (3)
C2—C3—C4—N2	-0.24 (19)	C12—C13—C14—S1	176.01 (14)
C2—C3—C4—C5	178.31 (17)	O2—S1—C14—C15	128.99 (15)
N2—C4—C5—C10	-133.78 (18)	O1—S1—C14—C15	0.77 (17)
C3—C4—C5—C10	47.9 (3)	N1—S1—C14—C15	-114.63 (15)
N2—C4—C5—C6	46.7 (2)	O2—S1—C14—C13	-48.55 (16)
C3—C4—C5—C6	-131.6 (2)	O1—S1—C14—C13	-176.76 (14)
C10—C5—C6—C7	1.8 (3)	N1—S1—C14—C13	67.84 (16)
C4—C5—C6—C7	-178.70 (16)	C13—C14—C15—C16	4.2 (3)
C5—C6—C7—C8	-0.3 (3)	S1—C14—C15—C16	-173.24 (13)
C6—C7—C8—C9	-1.4 (3)	C14—C15—C16—C11	-2.8 (3)
C7—C8—C9—C10	1.6 (3)	C12—C11—C16—C15	-1.2 (3)
C8—C9—C10—C5	-0.1 (3)	N2—C11—C16—C15	178.71 (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>
N1—H1...N3 <sup>i</sup>	0.92 (2)	1.98 (2)	2.878 (2)	164 (2)
N1—H2...O1 <sup>ii</sup>	0.86 (2)	2.07 (2)	2.930 (2)	177 (2)

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