organic compounds

Acta Crystallographica Section E **Structure Reports** Online

ISSN 1600-5368

3-Amino-5-bromo-2-iodopyridine

Kevin D. Bunker,^a Neal W. Sach,^a Seiji Nukui,^a Arnold L. Rheingold^b and Alex Yanovsky^a*

^aPfizer Global Research and Development, La Jolla Laboratories, 10614 Science Center Drive, San Diego, CA 92122, USA, and ^bDepartment of Chemistry and Biochemistry, University of California San Diego, 9500 Gilman Drive, La Jolla, CA 92093, USA

Correspondence e-mail: alex.yanovsky@pfizer.com

Received 27 November 2008; accepted 1 December 2008

Key indicators: single-crystal X-ray study; T = 100 K; mean σ (C–C) = 0.009 Å; R factor = 0.032; wR factor = 0.082; data-to-parameter ratio = 15.3.

The reaction of 3-amino-5-bromopyridine with N-iodosuccinimide in the presence of acetic acid produces the title compound, C₅H₄BrIN, with an iodo substituent in position 2 of the pyridine ring. The crystal structure features rather weak intermolecular N-H···N hydrogen bonds linking the molecules into chains along the z axis of the crystal.

Related literature

For structures of ortho-iodoanilines, see: McWilliam et al. (2001); Sandor & Foxman (2000); Parkin et al. (2005).



Experimental

Crystal data C₅H₄BrIN₂ $M_r = 298.90$

Monoclinic, $P2_1/c$ $a = 4.0983 (12) \text{ \AA}$

b = 15.172 (4) Å	Mo $K\alpha$ radiation
c = 12.038 (3) Å	$\mu = 9.53 \text{ mm}^{-1}$
$\beta = 90.152 \ (5)^{\circ}$	T = 100 (2) K
V = 748.5 (3) Å ³	$0.40 \times 0.33 \times 0.04 \text{ mm}$
$\mathbf{Z} - \mathbf{A}$	

Data collection

Refinement

1251 reflections

Bruker APEXII CCD	3783 measured reflections
diffractometer	1251 independent reflections
Absorption correction: multi-scan	1086 reflections with $I > 2\sigma(I)$
(SADABS; Bruker, 2001)	$R_{\rm int} = 0.037$
$T_{\min} = 0.234, T_{\max} = 0.557$	

 $\begin{array}{l} R[F^2>2\sigma(F^2)]=0.032\\ wR(F^2)=0.082 \end{array}$ 82 parameters H-atom parameters constrained $\Delta \rho_{\rm max} = 1.33 \ {\rm e} \ {\rm \AA}^{-3}$ $\Delta \rho_{\rm min} = -0.92 \text{ e} \text{ Å}^{-3}$

Table 1

S = 1.05

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	$D-{\rm H}$	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N2 - H2A \cdots N1^{i}$ $N2 - H2B \cdots I1$	0.88 0.88	2.16 2.79	3.025 (8) 3.259 (5)	166 115
Commentary and as (i) as				

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

Data collection: APEX2 (Bruker, 2007); cell refinement: SAINT (Bruker, 2007); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: RZ2275).

References

Bruker (2001). SADABS. Bruker AXS Inc., Madison, Wisconsin, USA. Bruker (2007). APEX2 and SAINT. Bruker AXS Inc., Madison, Wisconsin, USA.

McWilliam, S. A., Skakle, J. M. S., Low, J. N., Wardell, J. L., Garden, S. J., Pinto, A. C., Torres, J. C. & Glidewell, C. (2001). Acta Cryst. C57, 942-945.

Parkin, A., Spanswick, C. K., Pulham, C. R. & Wilson, C. C. (2005). Acta Cryst. E61 01087-01089

Sandor, R. B. & Foxman, B. M. (2000). Tetrahedron, 56, 6805-6812.

Sheldrick, G. M. (2008). Acta Cryst. A64, 112-122.

supplementary materials

Acta Cryst. (2009). E65, o28 [doi:10.1107/S1600536808040452]

3-Amino-5-bromo-2-iodopyridine

K. D. Bunker, N. W. Sach, S. Nukui, A. L. Rheingold and A. Yanovsky

Comment

The reaction of 5-bromo-3-aminopyridine with *N*-iodosuccinimide in the presence of acetic acid leads to iodo-substitution at position 2 of the pyridine ring, as shown by the X-ray study of the title compound (Fig. 1). To the best of our knowledge, this is the first structure of *ortho*-iodoaminopyridine derivative. The N2^{...}I1 distance 3.259 (5) Å is typical for *ortho*-iodoanilines (McWilliam *et al.*, 2001; Sandor & Foxman, 2000; Parkin *et al.*, 2005) and may suggest involvement of the H2B atom in weak intramolecular N2—H2B^{...}I1 interaction (Table 1).

The second `active' H-atom, H2A, participates in the intermolecular H-bond N2—H2A···N1ⁱ (symmetry code (i): x, 1/2 - y, z - 1/2; Table 1), which links the molecules into the chains along the z-axis of the crystal (Fig. 2). There are no strong halogen···halogen interactions in the structure; the shortest intermolecular I···I distances are 4.091 (1) Å and 4.098 (1) Å.

Experimental

To a solution of 3-amino-5-bromopyridine (100 mg, 0.56 mmol) in acetic acid (0.1 *M*, 5.61 ml) was added *N*-iodosuccinimide (133 mg, 0.56 mmol) at rt. After 3 h, the reaction was quenched with sat. sodium bicarbonate and extracted 3 times with EtOAc. The organic layers were combined, dried, filtered, and concentrated. The crude residue was subjected to flash chromatography (silica gel, 0–50% EtOAc/heptane). Isolated 93 mg (55%) of 3-amino-5-bromo-2-iodopyridine, as a brown solid. X-ray quality crystals were obtained by slow evaporation of a concentrated chromatography fraction (approx. 30% EtOAc/heptane). ¹H NMR (400 MHz, DMSO-d₆) (δ p.p.m.) 5.65 (s, 2 H), 7.16 (d, J = 2.27 Hz, 1 H), 7.67 (d, J = 2.01 Hz, 1 H). ¹³C NMR (101 MHz, DMSO-d₆) (δ p.p.m.) 106.10, 120.01, 120.92, 137.97, 147.68.

Refinement

All H atoms were treated as riding with the C—H and N—H distances of 0.95 Å and 0.88 Å respectively; the $U_{iso}(H)$ were set to $1.2U_{eq}$ of the carrying atom.

Figures



Fig. 1. Molecular structure of the title compound, showing 50% probability displacement ellipsoids and atom numbering scheme. H atoms are drawn as circles with arbitrary small radius.



Fig. 2. The crystal packing diagram viewed down the *x*-axis.

 $F_{000} = 544$

 $D_{\rm x} = 2.652 \ {\rm Mg \ m^{-3}}$ Mo Kα radiation

Cell parameters from 2536 reflections

 $\lambda = 0.71073 \text{ Å}$

 $\theta = 2.7 - 25.3^{\circ}$

3-Amino-5-bromo-2-iodopyridine

Crystal data C₅H₄BrIN₂ $M_r = 298.90$ Monoclinic, $P2_1/c$ Hall symbol: -P 2ybc a = 4.0983 (12) Å *b* = 15.172 (4) Å *c* = 12.038 (3) Å $\beta = 90.152 (5)^{\circ}$ V = 748.5 (3) Å³ Z = 4

Data collection

b = 15.172 (4) Å	$\mu = 9.53 \text{ mm}^{-1}$
c = 12.038 (3) Å	T = 100 (2) K
$\beta = 90.152 \ (5)^{\circ}$	Plate, colourless
V = 748.5 (3) Å ³	$0.40\times0.33\times0.04~mm$
Z = 4	
Data collection	
Bruker APEXII CCD diffractometer	1251 independent reflections
Radiation source: fine-focus sealed tube	1086 reflections with $I > 2\sigma(I)$
Monochromator: graphite	$R_{\rm int} = 0.037$
T = 100(2) K	$\theta_{\text{max}} = 25.3^{\circ}$

 $\theta_{\min} = 2.2^{\circ}$

 $h = -4 \rightarrow 1$ $k = -17 \rightarrow 18$ $l = -10 \rightarrow 14$

T = 100(2) K
ϕ and ω scans
Absorption correction: analytical
(SADABS; Bruker, 2001)
$T_{\min} = 0.234, T_{\max} = 0.557$
3783 measured reflections

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.032$	H-atom parameters constrained
$wR(F^2) = 0.082$	$w = 1/[\sigma^2(F_o^2) + (0.037P)^2 + 2.524P]$ where $P = (F_o^2 + 2F_c^2)/3$
<i>S</i> = 1.05	$(\Delta/\sigma)_{\rm max} = 0.003$
1251 reflections	$\Delta \rho_{max} = 1.33 \text{ e} \text{ Å}^{-3}$
82 parameters	$\Delta \rho_{min} = -0.92 \text{ e } \text{\AA}^{-3}$

Primary atom site location: structure-invariant direct 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 $F^2 > \sigma(F^2)$ 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.

	x	У	Ζ	$U_{\rm iso}$ */ $U_{\rm eq}$
I1	0.18405 (10)	0.37509 (3)	0.48761 (3)	0.02516 (17)
Br1	-0.56202 (16)	0.01782 (4)	0.28399 (5)	0.0266 (2)
N2	-0.0278 (15)	0.3306 (4)	0.2326 (4)	0.0298 (13)
H2A	-0.0808	0.3282	0.1617	0.036*
H2B	0.0760	0.3769	0.2589	0.036*
N1	-0.1133 (14)	0.1994 (4)	0.4858 (4)	0.0264 (13)
C2	-0.2684 (18)	0.1278 (4)	0.4470 (6)	0.0264 (15)
H2	-0.3243	0.0815	0.4965	0.032*
C1	-0.0390 (15)	0.2637 (5)	0.4165 (5)	0.0242 (14)
C5	-0.1071 (15)	0.2622 (5)	0.3019 (5)	0.0227 (14)
C3	-0.3480 (16)	0.1208 (4)	0.3350 (5)	0.0221 (14)
C4	-0.2713 (15)	0.1878 (4)	0.2636 (5)	0.0213 (14)
H4	-0.3303	0.1834	0.1874	0.026*

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\hat{A}^2)

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
I1	0.0244 (3)	0.0324 (3)	0.0187 (3)	-0.00094 (16)	-0.00522 (18)	-0.00340 (17)
Br1	0.0285 (4)	0.0274 (4)	0.0238 (4)	-0.0007 (3)	-0.0039 (3)	-0.0027 (3)
N2	0.044 (4)	0.029 (3)	0.016 (3)	-0.002 (3)	-0.008 (3)	-0.001 (2)
N1	0.035 (3)	0.030 (3)	0.015 (3)	0.002 (2)	-0.005 (2)	0.002 (2)
C2	0.035 (4)	0.024 (4)	0.020 (3)	0.000 (3)	-0.007 (3)	0.002 (3)
C1	0.017 (3)	0.036 (4)	0.020 (3)	0.004 (3)	-0.006 (3)	-0.008 (3)
C5	0.017 (3)	0.035 (4)	0.016 (3)	0.006 (3)	-0.001 (2)	-0.002 (3)
C3	0.021 (4)	0.027 (4)	0.018 (3)	0.005 (3)	-0.001 (3)	-0.001 (3)
C4	0.021 (3)	0.033 (4)	0.011 (3)	0.007 (3)	-0.005 (2)	-0.006 (3)
Geometric pa	rameters (Å, °)					
II—CI		2.102 (7)	C2—	-C3	1.390	0 (9)
Br1—C3		1.894 (7)	C2—	-H2	0.950	00

supplementary materials

N2—C5	1.371 (9)	C1—C5	1.407 (9)
N2—H2A	0.8800	C5—C4	1.393 (9)
N2—H2B	0.8800	C3—C4	1.368 (9)
N1—C1	1.320 (9)	C4—H4	0.9500
N1—C2	1.342 (9)		
C5—N2—H2A	120.0	N2C5C4	121.8 (5)
C5—N2—H2B	120.0	N2C5C1	122.6 (6)
H2A—N2—H2B	120.0	C4—C5—C1	115.6 (6)
C1—N1—C2	119.3 (5)	C4—C3—C2	119.9 (6)
N1—C2—C3	120.6 (6)	C4—C3—Br1	121.1 (5)
N1—C2—H2	119.7	C2—C3—Br1	119.0 (5)
С3—С2—Н2	119.7	C3—C4—C5	120.4 (6)
N1—C1—C5	124.2 (6)	С3—С4—Н4	119.8
N1—C1—I1	116.0 (4)	C5—C4—H4	119.8
C5—C1—I1	119.8 (5)		

Hydrogen-bond geometry (Å, °)

D—H···A	<i>D</i> —Н	$H \cdots A$	$D \cdots A$	$D\!\!-\!\!\mathrm{H}^{\ldots}\!A$
N2—H2A…N1 ⁱ	0.88	2.16	3.025 (8)	166
N2—H2B…I1	0.88	2.79	3.259 (5)	115
Symmetry codes: (i) x , $-y+1/2$, $z-1/2$.				





