

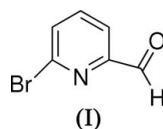
6-Bromopyridine-2-carbaldehyde

Huan-Xia Zhang,^a Da-Bin Qin,^{a*}
Lin-Hai Jing,^a Shao-Jin Gu^a and
Zhi-Hua Mao^b^aSchool of Chemistry and Chemical Industry,
China West Normal University, Nanchong
637002, People's Republic of China, and ^bThe
Centre of Test and Analysis, Sichuan University,
Chengdu 610064, People's Republic of ChinaCorrespondence e-mail:
zhanghuanxia@126.com

Key indicators

Single-crystal X-ray study
 $T = 290$ K
Mean $\sigma(\text{C}-\text{C}) = 0.007$ Å
 R factor = 0.038
 wR factor = 0.099
Data-to-parameter ratio = 12.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.The molecule of the title compound, $\text{C}_6\text{H}_4\text{BrNO}$, is essentially planar. In the crystal structure, $\text{C}-\text{H}\cdots\text{N}$ hydrogen bonds link the molecules into a chain along the b axis.Received 27 March 2006
Accepted 30 March 2006

Comment

6-Bromopyridine-2-carbaldehyde derivatives have received much attention as building blocks for supramolecules and ligands for transition metal catalysts and luminescent complexes (Li *et al.*, 2001; Orita *et al.*, 2004). We here report the crystal structure of the title compound, (I).Bond lengths and angles in (I) are normal (Table 1). The molecule is essentially planar, with an r.m.s. deviation of 0.006 (4) Å for fitted non-H atoms (Fig. 1). The crystal packing reveals that the molecules translated by one unit cell along the b -axis direction are linked into a chain by intermolecular $\text{C}-\text{H}\cdots\text{N}$ hydrogen-bonding interactions (Fig. 2).

Experimental

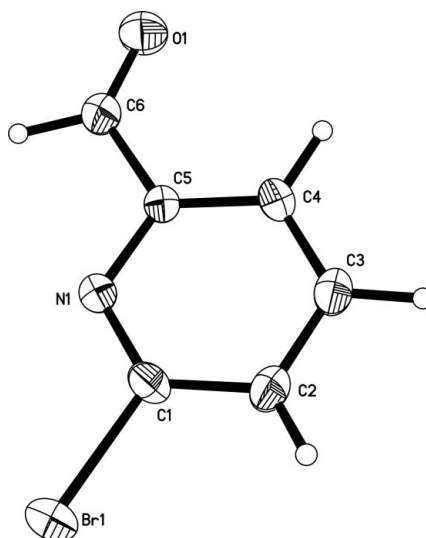
Compound (I) was prepared according to the reported procedure of Parks *et al.* (1971). Yellow single crystals suitable for X-ray diffraction were obtained by recrystallization from ethyl acetate.

Figure 1
The structure of (I), showing 30% probability displacement ellipsoids and the atomic numbering.

Crystal data

C₆H₄BrNO
M_r = 186.00
 Monoclinic, *P*₂₁/*a*
a = 6.908 (2) Å
b = 6.290 (4) Å
c = 15.060 (6) Å
 β = 95.57 (3)°
V = 651.3 (5) Å³

Z = 4
D_x = 1.897 Mg m⁻³
 Mo *K*α radiation
 μ = 6.22 mm⁻¹
T = 290 (2) K
 Block, yellow
 0.24 × 0.22 × 0.22 mm

Data collection

Enraf–Nonius CAD-4
 diffractometer
 ω/2θ scans
 Absorption correction: none
 1335 measured reflections
 1202 independent reflections

750 reflections with *I* > 2σ(*I*)
*R*_{int} = 0.012
 θ_{max} = 25.4°
 3 standard reflections
 every 200 reflections
 intensity decay: 1.3%

Refinement

Refinement on *F*²
R[*F*² > 2σ(*F*²)] = 0.038
wR(*F*²) = 0.099
S = 0.99
 1202 reflections
 99 parameters
 All H-atom parameters refined

w = 1/[σ²(*F*_o²) + (0.056*P*)²]
 where *P* = (*F*_o² + 2*F*_c²)/3
 (Δ/σ)_{max} = 0.001
 Δρ_{max} = 0.34 e Å⁻³
 Δρ_{min} = -0.44 e Å⁻³
 Extinction correction: *SHELXL97*
 Extinction coefficient: 0.037 (4)

Table 1

Selected geometric parameters (Å, °).

| | | | |
|-----------|-----------|----------|-----------|
| Br1—C1 | 1.901 (5) | C2—C3 | 1.368 (7) |
| O1—C6 | 1.207 (6) | C3—C4 | 1.364 (7) |
| N1—C1 | 1.296 (6) | C4—C5 | 1.376 (7) |
| N1—C5 | 1.341 (6) | C5—C6 | 1.481 (7) |
| C1—C2 | 1.390 (8) | | |
| C1—N1—C5 | 117.1 (4) | N1—C5—C6 | 114.8 (4) |
| N1—C1—C2 | 124.6 (4) | C4—C5—C6 | 122.5 (4) |
| N1—C1—Br1 | 116.4 (4) | O1—C6—C5 | 123.6 (5) |
| N1—C5—C4 | 122.7 (4) | | |

Table 2

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> |
|-------------------------|-------------|---------------|-----------------------|-------------------------|
| C3—H3...N1 ⁱ | 1.03 (6) | 2.52 (6) | 3.534 (7) | 167 (4) |

Symmetry code: (i) *x*, *y* + 1, *z*.

H atoms were located in a difference Fourier map and refined isotropically. The C—H bond lengths lie in the range 0.91 (5)–1.05 (6) Å.

Data collection: *DIFRAC* (Gabe *et al.*, 1993); cell refinement: *DIFRAC*; data reduction: *NRCVAX* (Gabe *et al.*, 1989); program(s)

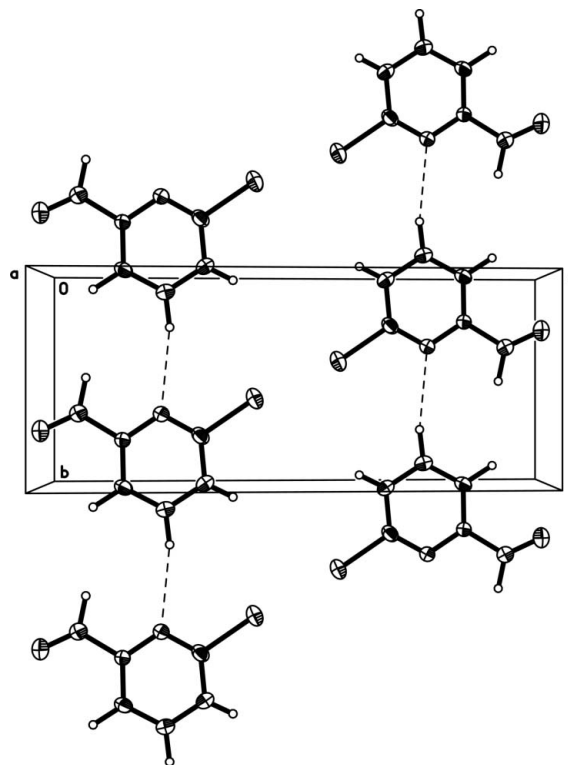


Figure 2

The crystal packing of (I), showing C—H...N hydrogen-bonded (dashed lines) chains.

used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *SHELXL97*.

The authors thank the Centre for Test and Analysis, Sichuan University, for financial support.

References

- Farrugia, L. J. (1997). *J. Appl. Cryst.* **30**, 565.
 Gabe, E. J., Le Page, Y., Charland, J.-P., Lee, F. L. & White, P. S. (1989). *J. Appl. Cryst.* **22**, 384–387.
 Gabe, E. J., White, P. S. & Enright, G. D. (1993). *DIFRAC*. National Research Council Canada, Ottawa.
 Li, X., Gibb, C. L. D., Kuebel, M. E. & Gibbp, B. C. (2001). *Tetrahedron*, **57**, 1175–1182.
 Orita, A., Nakano, T., Yokoyama, T., Babu, G. & Otera, J. (2004). *Chem. Lett.* **33**, 1298–1299.
 Parks, J. E., Wagner, B. E. & Holm, R. H. (1971). *Inorg. Chem.* **10**, 2472–2478.
 Sheldrick, G. M. (1997). *SHELXS97* and *SHELXL97*. University of Göttingen, Germany.