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Methyl 5-bromo-2-chloropyridine-3-carboxylate

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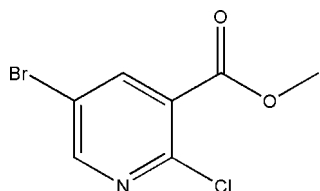
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.014$ Å; R factor = 0.045; wR factor = 0.131; data-to-parameter ratio = 8.4.

The title compound, $\text{C}_7\text{H}_5\text{BrClNO}_2$, crystallizes with two independent molecules in the asymmetric unit. In the absence of classical intermolecular interactions, the crystal structure exhibits relatively short intermolecular $\text{Br}\cdots\text{O}$ distances [3.143 (9) and 3.162 (9) Å].

Related literature

For the biological activity of the title compound, see: Colarusso & Narjes (2004); Kim *et al.* (2006). For related crystal structures, see McArdle *et al.* (1982).



Experimental

Crystal data

$\text{C}_7\text{H}_5\text{BrClNO}_2$
 $M_r = 250.48$

Triclinic, $P1$
 $a = 3.978$ (2) Å

$b = 8.153$ (3) Å
 $c = 14.040$ (2) Å
 $\alpha = 96.89$ (2)°
 $\beta = 96.20$ (3)°
 $\gamma = 100.70$ (2)°
 $V = 440.2$ (3) Å³

$Z = 2$
Mo $K\alpha$ radiation
 $\mu = 4.93$ mm⁻¹
 $T = 298$ (2) K
 $0.16 \times 0.14 \times 0.10$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2004)
 $T_{\min} = 0.506$, $T_{\max} = 0.638$

2186 measured reflections
1818 independent reflections
1564 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.130$
 $S = 1.09$
1818 reflections
217 parameters
3 restraints

H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 1.17$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.90$ e Å⁻³
Absolute structure: Flack (1983); 70 Friedel pairs
Flack parameter: 0.01 (2)

Table 1

Selected interatomic distances (Å).

$\text{Br1}\cdots\text{O3}^{\text{i}}$	3.143 (9)	$\text{Br2}\cdots\text{O1}^{\text{ii}}$	3.162 (9)
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Symmetry codes: (i) $x - 1, y - 1, z + 1$; (ii) $x - 1, y, z - 1$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINTE* (Bruker, 2001); data reduction: *SAINTE*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and local programs.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CV2402).

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supporting information

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Methyl 5-bromo-2-chloropyridine-3-carboxylate**Yi Ma and Ya-Lun Liu****S1. Comment**

The title compound, (I), is a useful intermediate for the synthesis of various bioactive compounds (Colarusso *et al.*, 2004; Kim *et al.*, 2006). In this paper, we report its crystal structure.

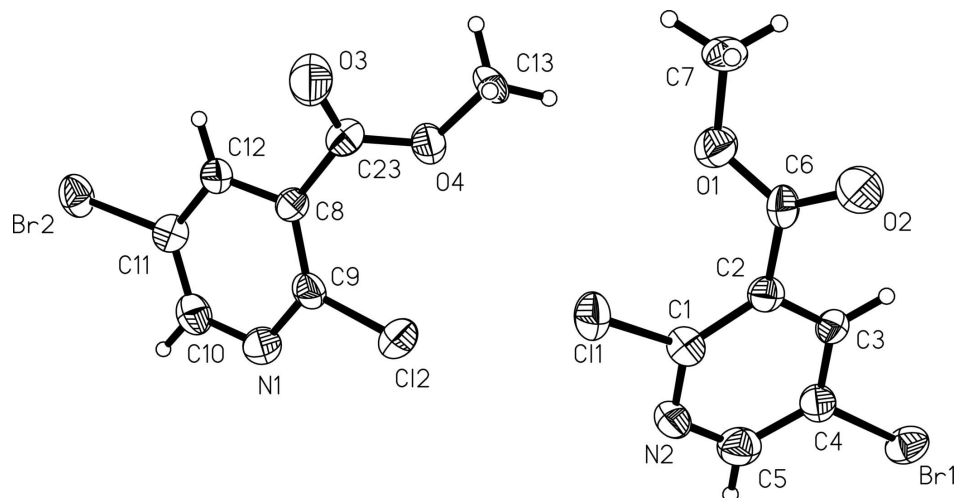
Compound (I) crystallizes with two independent molecules in the non-centrosymmetric triclinic unit cell (Fig. 1). The bond lengths and angles in the molecules are normal and in a good agreement with those reported previously (McArdle *et al.*, 1982). The dihedral angles between the planes of the methoxycarbonyl group (C6/C7/O1/O2; C13/C23/O3/O4) and pyridine rings in the two independent molecules are 45.8 (2) and 44.0 (3)°, respectively. In the absence of classical intermolecular interactions, the crystal packing exhibits relatively short intermolecular Br...O distances (Table 1).

S2. Experimental

A solution of 5-bromo-2-hydroxynicotinic acid (0.138 mol) and *N,N*-dimethylformamide (0.138 mol) in thionyl chloride (160 mL) was refluxed for 2 h. Thionyl chloride was evaporated and the yellow residue dissolved in anhydrous dichloromethane (200 mL), then anhydrous methanol was added dropwise. The resulting mixture was refluxed for 1 h and evaporated to afford slightly yellow oil which crystallized upon standing at room temperature. Crystals suitable for X-ray diffraction analysis were obtained by slow evaporation of an ethanol solution at room temperature over a period of one week.

S3. Refinement

All H atoms were placed in calculated positions, with C—H = 0.93 or 0.96 Å, and included in the final cycles of refinement using a riding model, with $U_{\text{iso}}(\text{H}) = 1.2\text{--}1.5$ times $U_{\text{eq}}(\text{C})$.

**Figure 1**

Two independent molecules of (I) with atomic numbering and displacement ellipsoids drawn at the 40% probability level.

Methyl 5-bromo-2-chloropyridine-3-carboxylate

Crystal data

$C_7H_5BrClNO_2$
 $M_r = 250.48$
 Triclinic, *P*1
 Hall symbol: P 1
 $a = 3.978$ (2) Å
 $b = 8.153$ (3) Å
 $c = 14.040$ (2) Å
 $\alpha = 96.89$ (2)°
 $\beta = 96.20$ (3)°
 $\gamma = 100.70$ (2)°
 $V = 440.2$ (3) Å³

$Z = 2$
 $F(000) = 244$
 $D_x = 1.890$ Mg m⁻³
 Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
 Cell parameters from 947 reflections
 $\theta = 2.6$ – 24.3 °
 $\mu = 4.93$ mm⁻¹
 $T = 298$ K
 Block, colourless
 $0.16 \times 0.14 \times 0.10$ mm

Data collection

Bruker SMART CCD area-detector
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 φ and ω scans
 Absorption correction: multi-scan
 (SADABS; Sheldrick, 2004)
 $T_{\min} = 0.506$, $T_{\max} = 0.639$

2186 measured reflections
 1818 independent reflections
 1564 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.017$
 $\theta_{\max} = 25.0$ °, $\theta_{\min} = 1.5$ °
 $h = -4 \rightarrow 3$
 $k = -9 \rightarrow 9$
 $l = -14 \rightarrow 16$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.130$
 $S = 1.09$
 1818 reflections
 217 parameters
 3 restraints

Primary atom site location: structure-invariant
 direct methods
 Secondary atom site location: difference Fourier
 map
 Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0806P)^2]$
 where $P = (F_o^2 + 2F_c^2)/3$

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 1.17 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.90 \text{ e } \text{\AA}^{-3}$$

Absolute structure: Flack (1983); 70 Friedel pairs
 Absolute structure parameter: 0.01 (2)

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.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	0.3498 (2)	0.81123 (11)	0.74398 (8)	0.0523 (3)
Br2	0.3533 (3)	1.31291 (12)	-0.24715 (8)	0.0530 (3)
Cl1	0.8539 (9)	0.7931 (4)	0.3331 (2)	0.0552 (7)
Cl2	1.2578 (8)	1.5050 (4)	0.1624 (2)	0.0555 (7)
O1	1.083 (3)	1.2390 (10)	0.5287 (6)	0.074 (3)
O2	0.762 (2)	1.1535 (9)	0.3865 (5)	0.0522 (19)
O3	1.295 (3)	1.8504 (10)	-0.0331 (6)	0.080 (3)
O4	1.116 (2)	1.8360 (9)	0.1109 (5)	0.0529 (19)
N1	0.579 (3)	0.6646 (10)	0.4701 (7)	0.050 (2)
N2	0.843 (3)	1.3036 (11)	0.0259 (7)	0.053 (2)
C1	0.703 (3)	0.8099 (12)	0.4449 (7)	0.040 (2)
C2	0.735 (3)	0.9674 (12)	0.5010 (7)	0.037 (2)
C3	0.627 (3)	0.9656 (12)	0.5894 (7)	0.038 (2)
H3A	0.6454	1.0663	0.6303	0.045*
C4	0.490 (3)	0.8134 (12)	0.6187 (7)	0.042 (2)
C5	0.464 (3)	0.6665 (14)	0.5574 (8)	0.055 (3)
H5A	0.3658	0.5650	0.5764	0.066*
C6	0.880 (3)	1.1312 (12)	0.4727 (7)	0.043 (2)
C7	0.885 (3)	1.3097 (13)	0.3515 (8)	0.050 (3)
H7A	0.7729	1.3062	0.2869	0.076*
H7B	1.1297	1.3255	0.3511	0.076*
H7C	0.8330	1.4015	0.3931	0.076*
C8	0.995 (3)	1.4644 (13)	0.0517 (8)	0.042 (2)
C9	0.973 (3)	1.5908 (12)	-0.0031 (7)	0.039 (2)
C10	0.783 (3)	1.5431 (12)	-0.0957 (7)	0.042 (2)
H10A	0.7706	1.6217	-0.1381	0.050*
C11	0.616 (3)	1.3777 (12)	-0.1223 (7)	0.040 (2)
C12	0.647 (3)	1.2631 (13)	-0.0611 (8)	0.049 (3)
H12A	0.5287	1.1523	-0.0800	0.059*
C13	1.144 (3)	1.7723 (13)	0.0234 (7)	0.042 (2)
C14	1.289 (3)	2.0069 (11)	0.1478 (8)	0.047 (3)
H14A	1.2448	2.0350	0.2130	0.071*

H14B	1.2055	2.0824	0.1079	0.071*
H14C	1.5334	2.0172	0.1470	0.071*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0550 (6)	0.0621 (6)	0.0479 (7)	0.0190 (5)	0.0166 (5)	0.0206 (5)
Br2	0.0487 (6)	0.0544 (6)	0.0496 (7)	0.0118 (4)	-0.0056 (5)	-0.0094 (5)
C11	0.068 (2)	0.0575 (15)	0.0389 (14)	0.0144 (14)	0.0110 (12)	-0.0028 (12)
C12	0.067 (2)	0.0589 (16)	0.0397 (14)	0.0138 (14)	-0.0012 (12)	0.0113 (12)
O1	0.092 (7)	0.049 (4)	0.064 (5)	-0.012 (5)	-0.013 (5)	0.010 (4)
O2	0.058 (5)	0.050 (4)	0.047 (4)	0.005 (4)	0.004 (4)	0.013 (3)
O3	0.115 (8)	0.056 (5)	0.059 (5)	-0.019 (5)	0.029 (5)	0.005 (4)
O4	0.073 (6)	0.041 (4)	0.042 (4)	0.005 (4)	0.014 (4)	-0.001 (3)
N1	0.059 (7)	0.034 (5)	0.053 (6)	0.009 (4)	-0.006 (5)	-0.001 (4)
N2	0.066 (7)	0.044 (5)	0.048 (6)	0.007 (5)	0.012 (5)	0.011 (4)
C1	0.035 (6)	0.041 (6)	0.043 (6)	0.006 (4)	0.007 (5)	0.003 (5)
C2	0.033 (6)	0.037 (5)	0.040 (5)	0.005 (4)	-0.002 (4)	0.006 (4)
C3	0.042 (6)	0.041 (5)	0.031 (5)	0.011 (4)	0.007 (4)	0.006 (4)
C4	0.039 (6)	0.045 (6)	0.043 (6)	0.013 (4)	0.002 (4)	0.004 (4)
C5	0.057 (8)	0.047 (6)	0.057 (7)	0.005 (5)	-0.001 (5)	0.010 (5)
C6	0.043 (7)	0.044 (6)	0.035 (5)	0.005 (5)	-0.001 (4)	-0.001 (4)
C7	0.070 (8)	0.037 (6)	0.049 (7)	0.005 (5)	0.024 (6)	0.019 (5)
C8	0.049 (7)	0.039 (6)	0.041 (6)	0.015 (5)	0.015 (5)	0.002 (4)
C9	0.047 (6)	0.038 (5)	0.033 (5)	0.006 (5)	0.013 (4)	0.006 (4)
C10	0.051 (7)	0.039 (5)	0.037 (5)	0.011 (5)	0.012 (4)	0.005 (4)
C11	0.031 (6)	0.045 (6)	0.044 (6)	0.004 (4)	0.004 (4)	0.005 (4)
C12	0.049 (7)	0.042 (6)	0.052 (6)	0.003 (5)	0.009 (5)	-0.001 (5)
C13	0.038 (6)	0.049 (6)	0.039 (6)	0.007 (5)	0.006 (4)	0.011 (4)
C14	0.060 (7)	0.024 (5)	0.051 (7)	0.003 (5)	0.000 (5)	-0.004 (4)

Geometric parameters (Å, °)

Br1—C4	1.902 (11)	C3—C4	1.389 (14)
Br2—C11	1.902 (10)	C3—H3A	0.9300
C11—C1	1.740 (10)	C4—C5	1.370 (15)
C12—C8	1.736 (12)	C5—H5A	0.9300
O1—C6	1.215 (11)	C7—H7A	0.9600
O2—C6	1.296 (11)	C7—H7B	0.9600
O2—C7	1.439 (12)	C7—H7C	0.9600
O3—C13	1.215 (11)	C8—C9	1.368 (14)
O4—C13	1.299 (13)	C9—C10	1.403 (13)
O4—C14	1.441 (12)	C9—C13	1.494 (14)
N1—C1	1.302 (12)	C10—C11	1.376 (13)
N1—C5	1.351 (15)	C10—H10A	0.9300
N2—C8	1.327 (13)	C11—C12	1.357 (13)
N2—C12	1.346 (14)	C12—H12A	0.9300
C1—C2	1.400 (13)	C14—H14A	0.9600

C2—C3	1.357 (13)	C14—H14B	0.9600
C2—C6	1.471 (13)	C14—H14C	0.9600
Br1...O3 ⁱ	3.143 (9)	Br2...O1 ⁱⁱ	3.162 (9)
C6—O2—C7	119.8 (8)	H7A—C7—H7C	109.5
C13—O4—C14	119.5 (9)	H7B—C7—H7C	109.5
C1—N1—C5	117.0 (9)	N2—C8—C9	125.6 (11)
C8—N2—C12	116.7 (9)	N2—C8—C12	114.0 (8)
N1—C1—C2	125.8 (9)	C9—C8—C12	120.4 (8)
N1—C1—C11	113.3 (8)	C8—C9—C10	116.4 (9)
C2—C1—C11	120.9 (7)	C8—C9—C13	126.8 (9)
C3—C2—C1	116.1 (8)	C10—C9—C13	116.7 (8)
C3—C2—C6	118.3 (9)	C11—C10—C9	118.6 (8)
C1—C2—C6	125.6 (9)	C11—C10—H10A	120.7
C2—C3—C4	120.0 (9)	C9—C10—H10A	120.7
C2—C3—H3A	120.0	C12—C11—C10	120.1 (9)
C4—C3—H3A	120.0	C12—C11—Br2	121.1 (7)
C5—C4—C3	119.2 (10)	C10—C11—Br2	118.8 (7)
C5—C4—Br1	121.0 (8)	N2—C12—C11	122.5 (9)
C3—C4—Br1	119.8 (7)	N2—C12—H12A	118.8
N1—C5—C4	122.0 (10)	C11—C12—H12A	118.8
N1—C5—H5A	119.0	O3—C13—O4	124.2 (9)
C4—C5—H5A	119.0	O3—C13—C9	121.9 (9)
O1—C6—O2	123.3 (9)	O4—C13—C9	114.0 (8)
O1—C6—C2	121.6 (9)	O4—C14—H14A	109.5
O2—C6—C2	115.0 (8)	O4—C14—H14B	109.5
O2—C7—H7A	109.5	H14A—C14—H14B	109.5
O2—C7—H7B	109.5	O4—C14—H14C	109.5
H7A—C7—H7B	109.5	H14A—C14—H14C	109.5
O2—C7—H7C	109.5	H14B—C14—H14C	109.5
C5—N1—C1—C2	1.3 (16)	C12—N2—C8—C9	-0.8 (17)
C5—N1—C1—C11	178.3 (8)	C12—N2—C8—C12	-177.4 (8)
N1—C1—C2—C3	0.5 (16)	N2—C8—C9—C10	-2.7 (16)
C11—C1—C2—C3	-176.3 (7)	C12—C8—C9—C10	173.7 (8)
N1—C1—C2—C6	179.1 (9)	N2—C8—C9—C13	-179.3 (11)
C11—C1—C2—C6	2.4 (15)	C12—C8—C9—C13	-3.0 (15)
C1—C2—C3—C4	-1.0 (14)	C8—C9—C10—C11	4.3 (14)
C6—C2—C3—C4	-179.8 (10)	C13—C9—C10—C11	-178.7 (9)
C2—C3—C4—C5	-0.1 (15)	C9—C10—C11—C12	-2.6 (15)
C2—C3—C4—Br1	178.4 (8)	C9—C10—C11—Br2	179.2 (8)
C1—N1—C5—C4	-2.5 (16)	C8—N2—C12—C11	2.8 (16)
C3—C4—C5—N1	2.0 (16)	C10—C11—C12—N2	-1.1 (16)
Br1—C4—C5—N1	-176.5 (8)	Br2—C11—C12—N2	177.1 (9)
C7—O2—C6—O1	2.9 (18)	C14—O4—C13—O3	-3.9 (17)
C7—O2—C6—C2	179.6 (10)	C14—O4—C13—C9	175.8 (10)
C3—C2—C6—O1	43.6 (16)	C8—C9—C13—O3	133.7 (12)

C1—C2—C6—O1	-135.1 (12)	C10—C9—C13—O3	-42.9 (15)
C3—C2—C6—O2	-133.2 (10)	C8—C9—C13—O4	-46.0 (15)
C1—C2—C6—O2	48.2 (15)	C10—C9—C13—O4	137.4 (10)

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