

4-Bromo-2-[(E)-(2-fluoro-5-nitrophenyl)-iminomethyl]phenol

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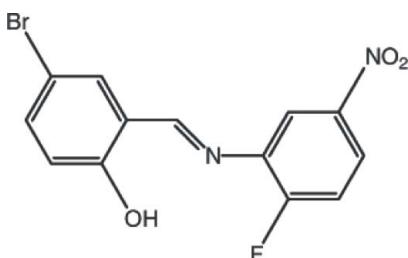
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.025; wR factor = 0.059; data-to-parameter ratio = 15.4.

The molecular conformation of the title compound, $\text{C}_{13}\text{H}_8\text{BrFN}_2\text{O}_3$, is essentially planar, with maximum deviations of 0.076 (1) and -0.080 (2) \AA for the O atoms of the NO_2 group. The molecular conformation is stabilized by an intramolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond, forming an $S(6)$ ring motif. In the crystal, pairs of molecules are linked via two pairs of $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds, forming inversion dimers that enclose $R_2^2(7)R_2^2(10)R_2^2(7)$ ring motifs.

Related literature

For the synthesis and biological activity of azomethines, see: Przybylski *et al.* (2009); Kalaivani *et al.* (2012); Blair *et al.* (2000). For the synthesis of fluorinated azomethines, see: Mohamed *et al.* (2012). For hydrogen-bond motifs, see: Bernstein *et al.* (1995). For standard bond lengths, see: Allen *et al.* (1987).



Experimental

Crystal data

$\text{C}_{13}\text{H}_8\text{BrFN}_2\text{O}_3$
 $M_r = 339.11$
Monoclinic, $P2_1/n$
 $a = 4.5082$ (9) \AA

$b = 19.815$ (4) \AA
 $c = 13.853$ (3) \AA
 $\beta = 95.484$ (5) $^\circ$
 $V = 1231.8$ (4) \AA^3

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 3.36\text{ mm}^{-1}$

$T = 100\text{ K}$
 $0.24 \times 0.04 \times 0.03\text{ mm}$

Data collection

Rigaku AFC12 (Right) diffractometer
Absorption correction: multi-scan (*CrystalClear-SM Expert*; Rigaku, 2012)
 $T_{\min} = 0.500$, $T_{\max} = 0.906$

8107 measured reflections
2811 independent reflections
2633 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.025$
 $wR(F^2) = 0.059$
 $S = 1.05$
2811 reflections

182 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.52\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.54\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1 \cdots N1	0.84	1.86	2.601 (2)	146
C7—H7 \cdots O3 ⁱ	0.95	2.45	3.399 (3)	173
C13—H13 \cdots O3 ⁱ	0.95	2.48	3.430 (3)	173

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

Data collection: *CrystalClear-SM Expert* (Rigaku, 2012); cell refinement: *CrystalClear-SM Expert*; data reduction: *CrystalClear-SM Expert*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012); software used to prepare material for publication: *WinGX* (Farrugia, 2012) and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5665).

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supporting information

Acta Cryst. (2013). E69, o107 [https://doi.org/10.1107/S1600536812050696]

4-Bromo-2-[(*E*)-(2-fluoro-5-nitrophenyl)iminomethyl]phenol

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S1. Comment

Schiff bases have been shown to exhibit a broad range of biological activities, including antifungal, antibacterial, antimalarial, antiproliferative, anti-inflammatory, antiviral, and antipyretic properties (Przybylski *et al.*, 2009; Kalaivani *et al.*, 2012). Among such compounds, the fluorinated Schiff's bases were considered to possess a distinguished biological activity due to the dramatic affect of fluorine atom on the metabolism and distribution of drug molecules in the body (Blair *et al.*, 2000). Further to our on going study on synthesis of bioactive fluorinated compounds (Mohamed *et al.*, 2012) we herein report the synthesis and crystal structure of a new fluorinated azomethine derivative.

In the title compound (I), (Fig. 1), the molecular conformation is essentially planar, with maximum deviations of 0.076 (1) and -0.080 (2) Å, respectively, for O2 and O3. The C1–C7–N1–C8 torsion angle is 179.92 (16)°. The bond lengths and angles in (I) are within the normal range (Allen *et al.*, 1987).

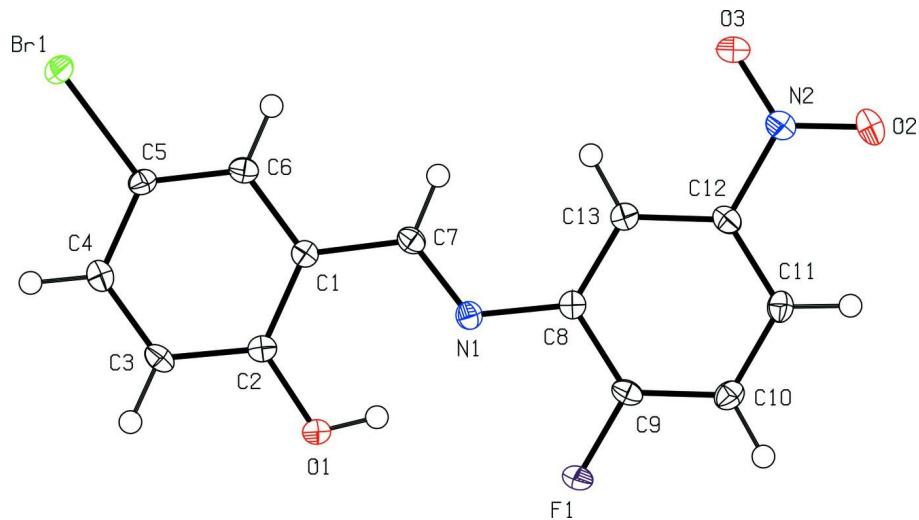
Molecular conformation is stabilized by O—H···N hydrogen bond (Table 1), forming an S(6) ring motif. In the crystal, the pairs of molecules are linked by C—H···O interactions (Table 1, Fig. 2), generating $R^2_2(7)R^2_2(10)R^2_2(7)$ ring motifs (Bernstein *et al.*, 1995) along the [001] direction.

S2. Experimental

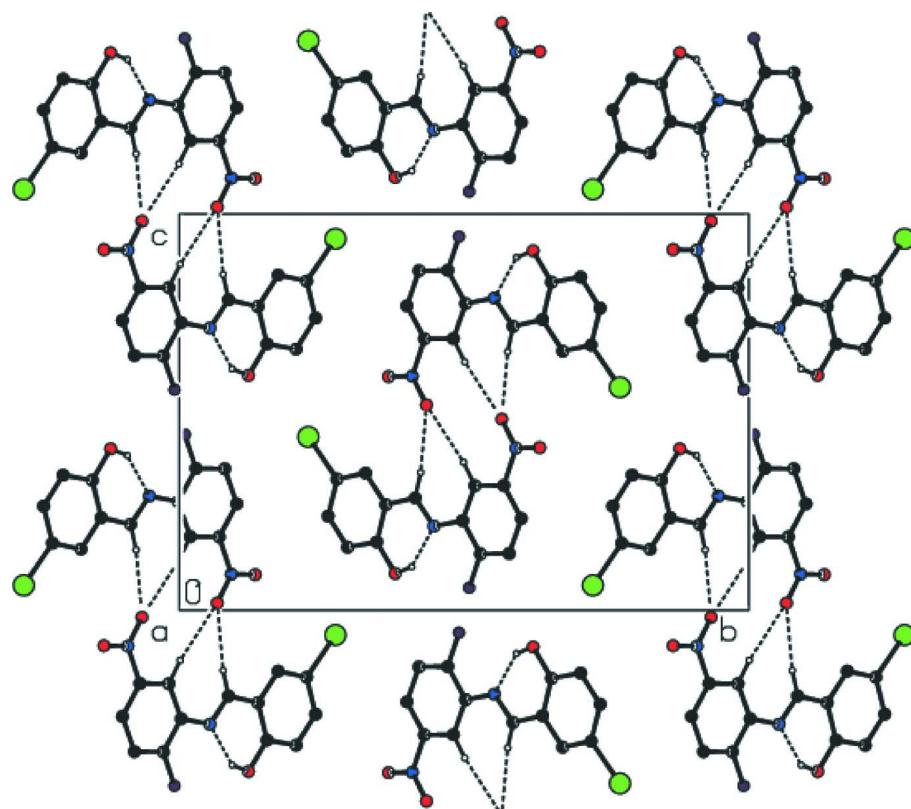
A mixture of 1 mmol (156 mg) 2-fluoro-5-nitroaniline and 1 mmol (201 mg) 5-bromo-2-hydroxybenzaldehyde in 50 ml ethanol was heated at 350 K and monitored by TLC till completion after 12 h. A mass solid product was deposited once the reaction mixture was allowed to cool at room temperature. The crude product was filtered dried under vacuum and washed by ethanol. Pure yellow rods (m.p. 465 K) suitable for X-ray diffraction were obtained in an excellent yield (92%) by crystallization of crude product from ethanol.

S3. Refinement

H atoms were positioned geometrically and refined using a riding model, with O—H = 0.84 Å, C—H = 0.95 Å, and with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{O})$ for hydroxyl and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ for the other H atoms.

**Figure 1**

The molecular structure of (I) with ellipsoids drawn at the 50% probability level.

**Figure 2**

Crystal packing of (I) viewed along the α axis. The hydrogen atoms not involved in the hydrogen bonds have been omitted for clarity.

4-Bromo-2-[(*E*)-(2-fluoro-5-nitrophenyl)iminomethyl]phenol*Crystal data*

$C_{13}H_8BrFN_2O_3$
 $M_r = 339.11$
Monoclinic, $P2_1/n$
Hall symbol: -P 2yn
 $a = 4.5082 (9)$ Å
 $b = 19.815 (4)$ Å
 $c = 13.853 (3)$ Å
 $\beta = 95.484 (5)^\circ$
 $V = 1231.8 (4)$ Å³
 $Z = 4$

$F(000) = 672$
 $D_x = 1.829$ Mg m⁻³
Mo $K\alpha$ radiation, $\lambda = 0.71075$ Å
Cell parameters from 4566 reflections
 $\theta = 2.5\text{--}31.2^\circ$
 $\mu = 3.36$ mm⁻¹
 $T = 100$ K
Rod, yellow
 $0.24 \times 0.04 \times 0.03$ mm

Data collection

Rigaku AFC12 (Right)
diffractometer
Radiation source: Rotating Anode
Detector resolution: 28.5714 pixels mm⁻¹
profile data from ω -scans
Absorption correction: multi-scan
(CrystalClear-SM Expert; Rigaku, 2012)
 $T_{\min} = 0.500$, $T_{\max} = 0.906$

8107 measured reflections
2811 independent reflections
2633 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.023$
 $\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 3.0^\circ$
 $h = -5 \rightarrow 5$
 $k = -25 \rightarrow 24$
 $l = -17 \rightarrow 14$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.025$
 $wR(F^2) = 0.059$
 $S = 1.05$
2811 reflections
182 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-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0255P)^2 + 1.1447P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.52$ e Å⁻³
 $\Delta\rho_{\min} = -0.54$ e Å⁻³

Special details

Geometry. Bond distances, angles etc. have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement on F^2 for ALL reflections except those flagged by the user for potential systematic errors. Weighted R -factors wR and all goodnesses 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 observed criterion of $F^2 > \sigma(F^2)$ is used only for calculating $-R$ -factor-obs 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 (Å²)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
Br1	-0.15532 (4)	0.22619 (1)	0.43736 (1)	0.0188 (1)
F1	0.7510 (2)	0.50903 (6)	0.05672 (7)	0.0202 (3)
O1	0.2171 (3)	0.37891 (7)	0.09274 (9)	0.0171 (4)
O2	1.5039 (3)	0.63282 (8)	0.41030 (11)	0.0290 (4)

O3	1.2227 (5)	0.56614 (11)	0.48212 (12)	0.0614 (8)
N1	0.5847 (3)	0.44689 (7)	0.21201 (11)	0.0140 (4)
N2	1.3057 (4)	0.59101 (9)	0.40914 (12)	0.0227 (5)
C1	0.2662 (4)	0.36222 (8)	0.26646 (12)	0.0123 (5)
C2	0.1398 (4)	0.34652 (9)	0.17204 (12)	0.0135 (5)
C3	-0.0742 (4)	0.29535 (9)	0.15895 (13)	0.0157 (5)
C4	-0.1635 (4)	0.26099 (9)	0.23746 (13)	0.0159 (5)
C5	-0.0389 (4)	0.27669 (9)	0.33084 (12)	0.0140 (5)
C6	0.1728 (4)	0.32683 (9)	0.34603 (12)	0.0145 (5)
C7	0.4936 (4)	0.41399 (9)	0.28338 (13)	0.0138 (5)
C8	0.8050 (4)	0.49730 (9)	0.22592 (13)	0.0136 (5)
C9	0.8877 (4)	0.52873 (9)	0.14215 (12)	0.0150 (5)
C10	1.0992 (4)	0.57931 (9)	0.14290 (13)	0.0165 (5)
C11	1.2365 (4)	0.60049 (9)	0.23159 (13)	0.0157 (5)
C12	1.1566 (4)	0.56969 (9)	0.31484 (13)	0.0156 (5)
C13	0.9451 (4)	0.51880 (9)	0.31435 (13)	0.0154 (5)
H1	0.34830	0.40790	0.10930	0.0260*
H3	-0.15830	0.28430	0.09540	0.0190*
H4	-0.31010	0.22660	0.22810	0.0190*
H6	0.25500	0.33730	0.41000	0.0170*
H7	0.57530	0.42350	0.34770	0.0170*
H10	1.14920	0.59910	0.08410	0.0200*
H11	1.38190	0.63530	0.23510	0.0190*
H13	0.89690	0.49900	0.37330	0.0180*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Br1	0.0189 (1)	0.0213 (1)	0.0161 (1)	-0.0045 (1)	0.0003 (1)	0.0053 (1)
F1	0.0209 (6)	0.0273 (6)	0.0115 (5)	-0.0064 (5)	-0.0028 (4)	0.0011 (4)
O1	0.0208 (7)	0.0184 (6)	0.0119 (6)	-0.0042 (5)	0.0007 (5)	0.0008 (5)
O2	0.0326 (8)	0.0292 (8)	0.0243 (7)	-0.0159 (7)	-0.0022 (6)	-0.0043 (6)
O3	0.0880 (16)	0.0804 (15)	0.0135 (8)	-0.0628 (13)	-0.0074 (9)	0.0082 (8)
N1	0.0128 (7)	0.0141 (7)	0.0150 (7)	0.0001 (6)	0.0013 (6)	-0.0005 (5)
N2	0.0290 (9)	0.0228 (8)	0.0156 (8)	-0.0085 (7)	-0.0011 (7)	-0.0008 (6)
C1	0.0110 (8)	0.0124 (8)	0.0136 (8)	0.0021 (6)	0.0011 (6)	-0.0003 (6)
C2	0.0137 (8)	0.0135 (8)	0.0133 (8)	0.0026 (6)	0.0015 (6)	0.0004 (6)
C3	0.0156 (9)	0.0171 (8)	0.0140 (8)	0.0000 (7)	-0.0011 (6)	-0.0047 (7)
C4	0.0145 (8)	0.0144 (8)	0.0186 (9)	-0.0005 (7)	0.0001 (7)	-0.0018 (7)
C5	0.0144 (8)	0.0148 (8)	0.0127 (8)	0.0009 (7)	0.0015 (6)	0.0021 (6)
C6	0.0147 (8)	0.0160 (8)	0.0125 (8)	0.0014 (7)	0.0000 (6)	0.0002 (6)
C7	0.0123 (8)	0.0151 (8)	0.0137 (8)	0.0012 (7)	-0.0005 (6)	-0.0014 (6)
C8	0.0129 (8)	0.0129 (8)	0.0151 (8)	0.0007 (6)	0.0016 (7)	-0.0003 (6)
C9	0.0139 (8)	0.0179 (8)	0.0125 (8)	0.0022 (7)	-0.0023 (6)	-0.0014 (6)
C10	0.0164 (9)	0.0166 (8)	0.0167 (9)	0.0011 (7)	0.0022 (7)	0.0036 (7)
C11	0.0155 (8)	0.0126 (8)	0.0191 (9)	-0.0009 (7)	0.0017 (7)	0.0007 (7)
C12	0.0163 (9)	0.0155 (8)	0.0145 (8)	-0.0004 (7)	-0.0010 (7)	-0.0021 (6)
C13	0.0160 (9)	0.0159 (8)	0.0143 (8)	-0.0009 (7)	0.0012 (7)	0.0007 (6)

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

Br1—C5	1.8980 (18)	C5—C6	1.380 (3)
F1—C9	1.339 (2)	C8—C13	1.390 (3)
O1—C2	1.347 (2)	C8—C9	1.399 (2)
O2—N2	1.218 (2)	C9—C10	1.383 (3)
O3—N2	1.215 (3)	C10—C11	1.387 (3)
O1—H1	0.8400	C11—C12	1.383 (3)
N1—C7	1.284 (2)	C12—C13	1.387 (3)
N1—C8	1.409 (2)	C3—H3	0.9500
N2—C12	1.473 (2)	C4—H4	0.9500
C1—C6	1.405 (2)	C6—H6	0.9500
C1—C7	1.453 (2)	C7—H7	0.9500
C1—C2	1.411 (2)	C10—H10	0.9500
C2—C3	1.399 (3)	C11—H11	0.9500
C3—C4	1.376 (3)	C13—H13	0.9500
C4—C5	1.395 (2)		
C2—O1—H1	109.00	F1—C9—C10	118.47 (15)
C7—N1—C8	121.86 (16)	C8—C9—C10	123.70 (16)
O2—N2—C12	118.63 (16)	C9—C10—C11	118.36 (16)
O3—N2—C12	118.09 (18)	C10—C11—C12	118.40 (17)
O2—N2—O3	123.29 (18)	N2—C12—C11	118.69 (16)
C2—C1—C7	121.43 (15)	C11—C12—C13	123.40 (17)
C6—C1—C7	119.05 (15)	N2—C12—C13	117.91 (16)
C2—C1—C6	119.52 (16)	C8—C13—C12	118.76 (16)
O1—C2—C3	117.94 (15)	C2—C3—H3	120.00
C1—C2—C3	119.50 (16)	C4—C3—H3	120.00
O1—C2—C1	122.56 (16)	C3—C4—H4	120.00
C2—C3—C4	120.43 (16)	C5—C4—H4	120.00
C3—C4—C5	120.02 (17)	C1—C6—H6	120.00
Br1—C5—C4	119.11 (13)	C5—C6—H6	120.00
Br1—C5—C6	119.98 (13)	N1—C7—H7	120.00
C4—C5—C6	120.88 (16)	C1—C7—H7	120.00
C1—C6—C5	119.65 (15)	C9—C10—H10	121.00
N1—C7—C1	120.45 (16)	C11—C10—H10	121.00
N1—C8—C9	116.27 (16)	C10—C11—H11	121.00
N1—C8—C13	126.35 (16)	C12—C11—H11	121.00
C9—C8—C13	117.38 (16)	C8—C13—H13	121.00
F1—C9—C8	117.83 (16)	C12—C13—H13	121.00
C8—N1—C7—C1	-179.92 (16)	C3—C4—C5—C6	0.4 (3)
C7—N1—C8—C9	179.13 (17)	C3—C4—C5—Br1	-177.66 (14)
C7—N1—C8—C13	-1.3 (3)	Br1—C5—C6—C1	177.55 (13)
O3—N2—C12—C13	-4.7 (3)	C4—C5—C6—C1	-0.5 (3)
O2—N2—C12—C11	-3.3 (3)	N1—C8—C9—C10	179.72 (16)
O2—N2—C12—C13	175.80 (17)	C13—C8—C9—F1	-179.08 (16)
O3—N2—C12—C11	176.20 (19)	C13—C8—C9—C10	0.1 (3)

C7—C1—C2—C3	178.71 (17)	N1—C8—C9—F1	0.5 (2)
C7—C1—C6—C5	−178.80 (17)	N1—C8—C13—C12	−179.49 (17)
C2—C1—C7—N1	0.1 (3)	C9—C8—C13—C12	0.1 (3)
C2—C1—C6—C5	0.7 (3)	F1—C9—C10—C11	178.83 (16)
C6—C1—C2—C3	−0.8 (3)	C8—C9—C10—C11	−0.4 (3)
C7—C1—C2—O1	−1.2 (3)	C9—C10—C11—C12	0.4 (3)
C6—C1—C7—N1	179.58 (16)	C10—C11—C12—N2	178.83 (16)
C6—C1—C2—O1	179.35 (16)	C10—C11—C12—C13	−0.3 (3)
O1—C2—C3—C4	−179.43 (16)	N2—C12—C13—C8	−179.08 (16)
C1—C2—C3—C4	0.7 (3)	C11—C12—C13—C8	0.0 (3)
C2—C3—C4—C5	−0.5 (3)		

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
O1—H1···N1	0.84	1.86	2.601 (2)	146
C7—H7···O3 ⁱ	0.95	2.45	3.399 (3)	173
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Symmetry code: (i) $-x+2, -y+1, -z+1$.