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3-Fluoro-4-(4-hydroxyphenoxy)-benzonitrile

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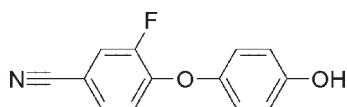
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Key indicators: single-crystal X-ray study; $T = 295$ K; mean $\sigma(\text{C}-\text{C}) = 0.005$ Å; R factor = 0.037; wR factor = 0.117; data-to-parameter ratio = 9.6.

The title compound, $\text{C}_{13}\text{H}_8\text{FNO}_2$, was synthesized from 3,4-difluorobenzonitrile and hydroquinone. The dihedral angle between the two aromatic rings is $70.9(2)^\circ$. In the crystal structure, molecules are linked by $\text{O}-\text{H}\cdots\text{N}$ hydrogen bonds, forming zigzag chains.

Related literature

For the herbicidal activity of hydroquinone derivatives, see: Bao *et al.* (2007); Liu (2002). For related structures, see: Sørensen *et al.* (2009); Luo *et al.* (2009); Zhang *et al.* (2009).



Experimental

Crystal data

 $\text{C}_{13}\text{H}_8\text{FNO}_2$ $M_r = 229.20$ Orthorhombic, $P2_12_12_1$ $a = 6.1932(4)$ Å $b = 8.8109(5)$ Å $c = 20.5269(12)$ Å $V = 1120.11(12)$ Å³ $Z = 4$ Mo $K\alpha$ radiation $\mu = 0.10$ mm⁻¹ $T = 295$ K $0.39 \times 0.31 \times 0.22$ mm

Data collection

Rigaku R-Axis RAPID diffractometer
Absorption correction: multi-scan (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.959$, $T_{\max} = 0.976$

10999 measured reflections
1498 independent reflections
928 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.032$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$
 $wR(F^2) = 0.117$
 $S = 1.01$
1498 reflections

156 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.17$ e Å⁻³
 $\Delta\rho_{\min} = -0.15$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O2}-\text{H201}\cdots\text{N1}^i$	0.82	2.03	2.839 (4)	168

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

Data collection: *PROCESS-AUTO* (Rigaku, 2006); cell refinement: *PROCESS-AUTO*; data reduction: *CrystalStructure* (Rigaku/MS, 2007); 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, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

The authors are grateful to Mr Jianming Gu for the crystal structure analysis.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: IM2202).

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supporting information

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3-Fluoro-4-(4-hydroxyphenoxy)benzotrile

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S1. Comment

Hydroquinone derivatives are important intermediates of herbicide synthesis and have therefore received growing attention recently (Liu, 2002; Bao *et al.*, 2007). Several hydroquinone derivatives were synthesized and investigated by X-ray diffraction in our laboratory. 4-(4-Cyano-2-fluoro-phenoxy)-phenol was obtained reacting hydroquinone and 3,4-difluorobenzotrile and its molecular structure is shown in Fig.1.

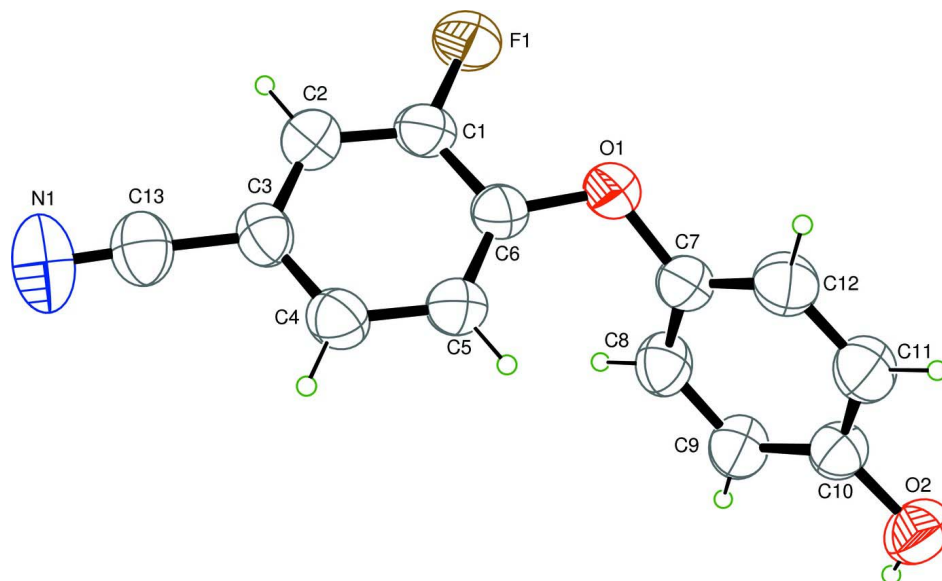
As it is expected substituents at both aromatic rings are coplanar with respect to the aromatic planes. The dihedral angle between the two planes is 70.66°. The molecule is bent with a C6—O1—C7 angle of 118.0 (2)°. The crystal structure is determined by intermolecular O—H···N interactions. The resulting supramolecular chains of the title compound showing H-bridge interactions is shown in Fig.2.

S2. Experimental

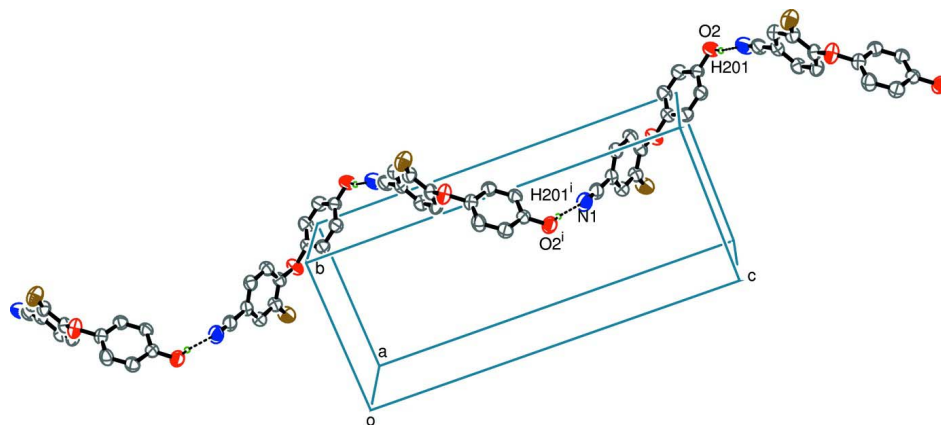
A DMSO (10 ml) solution of hydroquinone (0.0012 mol) and NaOH (0.0024 mol) was stirred at room temperature for 5 h. Then the mixture was heated to 80°C and 3,4-difluorobenzotrile (0.001 mol) was added dropwise and stirred for 10 h. Then the mixture was washed with water (30 ml) and extracted with ethyl acetate (three times). The organic solvent was removed under reduced pressure and the resulting crude product was purified by silica gel chromatography (pentane: ethyl acetate mixtures, yield 86%). Single crystals were obtained by slow evaporation of ethyl acetate at room temperature.

S3. Refinement

In the absence of significant anomalous dispersion effects, Friedel pairs were averaged. H atoms were placed in calculated positions with C—H = 0.98 Å (sp), C—H = 0.97 Å (sp²), C—H = 0.93 Å (aromatic). All H atoms were included in the final cycles of refinement using a riding model, with $U_{\text{iso}}(\text{H})=1.2U_{\text{eq}}$ of the respective carrier atoms.

**Figure 1**

Molecular structure of title compound, with the atomic labeling scheme. Displacement ellipsoids are drawn at the 50% probability level.

**Figure 2**

A partial packing diagram of title compound. Hydrogen bonds are shown as dashed lines. [Symmetry code: (i) $-x+1/2, -y+2, z+1/2$].

3-Fluoro-4-(4-hydroxyphenoxy)benzonitrile

Crystal data

$C_{13}H_8FNO_2$

$M_r = 229.20$

Orthorhombic, $P2_12_12_1$

Hall symbol: $P\ 2ac\ 2ab$

$a = 6.1932\ (4)\ \text{\AA}$

$b = 8.8109\ (5)\ \text{\AA}$

$c = 20.5269\ (12)\ \text{\AA}$

$V = 1120.11\ (12)\ \text{\AA}^3$

$Z = 4$

$F(000) = 472$

$D_x = 1.359\ \text{Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073\ \text{\AA}$

Cell parameters from 6842 reflections

$\theta = 3.0\text{--}27.4^\circ$

$\mu = 0.10\ \text{mm}^{-1}$

$T = 295\ \text{K}$

Chunk, colorless

$0.39 \times 0.31 \times 0.22\ \text{mm}$

Data collection

Rigaku R-AXIS RAPID diffractometer	10999 measured reflections 1498 independent reflections
Radiation source: rolling anode	928 reflections with $I > 2\sigma(I)$
Graphite monochromator	$R_{\text{int}} = 0.032$
Detector resolution: 10.00 pixels mm^{-1}	$\theta_{\text{max}} = 27.4^\circ$, $\theta_{\text{min}} = 3.1^\circ$
ω scans	$h = -8 \rightarrow 8$
Absorption correction: multi-scan (<i>ABSCOR</i> ; Higashi, 1995)	$k = -11 \rightarrow 11$
$T_{\text{min}} = 0.959$, $T_{\text{max}} = 0.976$	$l = -26 \rightarrow 26$

Refinement

Refinement on F^2	Hydrogen site location: inferred from neighbouring sites
Least-squares matrix: full	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.037$	$w = 1/[\sigma^2(F_o^2) + (0.0487P)^2 + 0.2503P]$
$wR(F^2) = 0.117$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.01$	$(\Delta/\sigma)_{\text{max}} < 0.001$
1498 reflections	$\Delta\rho_{\text{max}} = 0.17 \text{ e } \text{\AA}^{-3}$
156 parameters	$\Delta\rho_{\text{min}} = -0.15 \text{ e } \text{\AA}^{-3}$
0 restraints	Extinction correction: <i>SHELXL97</i> (Sheldrick, 2008), $F_c^* = kFc[1 + 0.001x\text{Fc}^2\lambda^3/\sin(2\theta)]^{-1/4}$
Primary atom site location: structure-invariant direct methods	Extinction coefficient: 0.031 (5)
Secondary atom site location: difference Fourier map	

Special details

Geometry. All esds (except the esd in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell esds are taken into account individually in the estimation of esds in distances, angles and torsion angles; correlations between esds in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell esds is used for estimating esds 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)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.8260 (4)	0.8262 (3)	0.91021 (10)	0.0842 (7)
F1	0.7477 (4)	0.5751 (2)	0.84401 (10)	0.0994 (7)
C3	0.2965 (5)	0.7967 (4)	0.78841 (13)	0.0647 (8)
C13	0.1155 (6)	0.7849 (4)	0.74589 (16)	0.0797 (10)
C7	0.8277 (5)	0.9255 (4)	0.96363 (13)	0.0662 (8)
O2	0.8846 (4)	1.2010 (3)	1.12504 (11)	0.0890 (8)
H201	0.7775	1.1944	1.1484	0.134*
C4	0.3341 (5)	0.9302 (4)	0.82157 (14)	0.0706 (8)
H4	0.2411	1.0120	0.8158	0.085*
C10	0.8605 (5)	1.1076 (3)	1.07184 (13)	0.0647 (8)
C1	0.6082 (5)	0.6918 (4)	0.83712 (14)	0.0689 (8)
C8	0.6707 (5)	0.9204 (4)	1.01045 (14)	0.0720 (8)
H8	0.5543	0.8546	1.0059	0.086*
C6	0.6460 (5)	0.8233 (4)	0.87182 (13)	0.0648 (8)

C2	0.4369 (5)	0.6748 (4)	0.79642 (14)	0.0712 (8)
H2	0.4141	0.5839	0.7744	0.085*
C9	0.6853 (5)	1.0130 (4)	1.06440 (14)	0.0718 (9)
H9	0.5769	1.0115	1.0958	0.086*
C11	1.0188 (6)	1.1111 (4)	1.02514 (15)	0.0789 (10)
H11	1.1376	1.1745	1.0301	0.095*
C12	1.0011 (6)	1.0202 (4)	0.97087 (15)	0.0805 (10)
H12	1.1076	1.0231	0.9390	0.097*
C5	0.5080 (6)	0.9438 (4)	0.86314 (14)	0.0710 (9)
H5	0.5319	1.0344	0.8853	0.085*
N1	-0.0310 (6)	0.7800 (4)	0.71268 (15)	0.1078 (12)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0678 (14)	0.1042 (16)	0.0805 (13)	0.0258 (14)	-0.0173 (12)	-0.0286 (13)
F1	0.0986 (16)	0.0867 (12)	0.1129 (14)	0.0398 (13)	-0.0214 (12)	-0.0218 (12)
C3	0.0581 (18)	0.083 (2)	0.0534 (14)	-0.0009 (17)	-0.0006 (13)	0.0069 (16)
C13	0.076 (2)	0.092 (2)	0.0710 (19)	-0.009 (2)	-0.0057 (19)	0.0122 (18)
C7	0.0610 (18)	0.0736 (17)	0.0642 (15)	0.0089 (17)	-0.0031 (16)	-0.0066 (15)
O2	0.0893 (19)	0.0916 (15)	0.0861 (15)	-0.0181 (15)	0.0185 (13)	-0.0244 (14)
C4	0.069 (2)	0.0754 (19)	0.0671 (16)	0.0133 (18)	-0.0032 (17)	0.0033 (16)
C10	0.066 (2)	0.0648 (17)	0.0632 (15)	-0.0042 (16)	0.0052 (16)	-0.0025 (14)
C1	0.068 (2)	0.0692 (18)	0.0697 (17)	0.0159 (17)	0.0009 (16)	-0.0051 (17)
C8	0.065 (2)	0.0776 (18)	0.0733 (17)	-0.0135 (18)	0.0016 (17)	-0.0006 (17)
C6	0.0592 (19)	0.0778 (18)	0.0572 (15)	0.0090 (17)	0.0002 (14)	-0.0069 (15)
C2	0.072 (2)	0.0751 (19)	0.0665 (17)	0.0011 (18)	-0.0016 (16)	-0.0044 (17)
C9	0.068 (2)	0.084 (2)	0.0632 (16)	-0.0165 (19)	0.0125 (16)	-0.0001 (16)
C11	0.063 (2)	0.094 (2)	0.0801 (19)	-0.0177 (19)	0.0167 (18)	-0.0068 (19)
C12	0.066 (2)	0.107 (2)	0.0689 (17)	-0.004 (2)	0.0148 (18)	-0.0094 (19)
C5	0.072 (2)	0.0706 (18)	0.0706 (17)	0.0124 (17)	-0.0058 (17)	-0.0094 (17)
N1	0.094 (2)	0.128 (3)	0.101 (2)	-0.030 (2)	-0.031 (2)	0.034 (2)

Geometric parameters (Å, °)

O1—C6	1.365 (4)	C10—C11	1.371 (4)
O1—C7	1.403 (4)	C10—C9	1.377 (4)
F1—C1	1.350 (3)	C1—C2	1.359 (4)
C3—C4	1.379 (5)	C1—C6	1.380 (4)
C3—C2	1.391 (4)	C8—C9	1.378 (4)
C3—C13	1.425 (4)	C8—H8	0.9300
C13—N1	1.135 (4)	C6—C5	1.375 (4)
C7—C12	1.368 (5)	C2—H2	0.9300
C7—C8	1.368 (4)	C9—H9	0.9300
O2—C10	1.375 (3)	C11—C12	1.376 (4)
O2—H201	0.8200	C11—H11	0.9300
C4—C5	1.379 (4)	C12—H12	0.9300
C4—H4	0.9300	C5—H5	0.9300

C6—O1—C7	118.0 (2)	C9—C8—H8	120.0
C4—C3—C2	119.7 (3)	O1—C6—C5	124.6 (3)
C4—C3—C13	119.8 (3)	O1—C6—C1	116.9 (3)
C2—C3—C13	120.5 (3)	C5—C6—C1	118.4 (3)
N1—C13—C3	177.8 (5)	C1—C2—C3	118.4 (3)
C12—C7—C8	120.1 (3)	C1—C2—H2	120.8
C12—C7—O1	118.1 (3)	C3—C2—H2	120.8
C8—C7—O1	121.6 (3)	C10—C9—C8	119.9 (3)
C10—O2—H201	109.5	C10—C9—H9	120.0
C5—C4—C3	120.7 (3)	C8—C9—H9	120.0
C5—C4—H4	119.6	C10—C11—C12	119.7 (3)
C3—C4—H4	119.6	C10—C11—H11	120.1
C11—C10—O2	117.7 (3)	C12—C11—H11	120.1
C11—C10—C9	119.9 (3)	C7—C12—C11	120.3 (3)
O2—C10—C9	122.4 (3)	C7—C12—H12	119.8
F1—C1—C2	118.7 (3)	C11—C12—H12	119.8
F1—C1—C6	118.5 (3)	C6—C5—C4	119.9 (3)
C2—C1—C6	122.8 (3)	C6—C5—H5	120.0
C7—C8—C9	119.9 (3)	C4—C5—H5	120.0
C7—C8—H8	120.0		
C6—O1—C7—C12	-129.3 (3)	C4—C3—C2—C1	0.0 (5)
C6—O1—C7—C8	56.0 (4)	C13—C3—C2—C1	179.5 (3)
C2—C3—C4—C5	-0.7 (5)	C11—C10—C9—C8	0.8 (5)
C13—C3—C4—C5	179.8 (3)	O2—C10—C9—C8	-179.2 (3)
C12—C7—C8—C9	1.2 (5)	C7—C8—C9—C10	-1.6 (5)
O1—C7—C8—C9	175.8 (3)	O2—C10—C11—C12	-179.7 (3)
C7—O1—C6—C5	27.9 (4)	C9—C10—C11—C12	0.3 (5)
C7—O1—C6—C1	-155.4 (3)	C8—C7—C12—C11	-0.1 (5)
F1—C1—C6—O1	1.0 (4)	O1—C7—C12—C11	-174.8 (3)
C2—C1—C6—O1	-179.0 (3)	C10—C11—C12—C7	-0.7 (5)
F1—C1—C6—C5	177.9 (3)	O1—C6—C5—C4	178.0 (3)
C2—C1—C6—C5	-2.1 (5)	C1—C6—C5—C4	1.4 (5)
F1—C1—C2—C3	-178.6 (3)	C3—C4—C5—C6	0.0 (5)
C6—C1—C2—C3	1.4 (5)		

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>
O2—H201...N1 ⁱ	0.82	2.03	2.839 (4)	168

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