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(E)-4-[(5-Methyl-2-furyl)methylene-amino]benzenesulfonic acid

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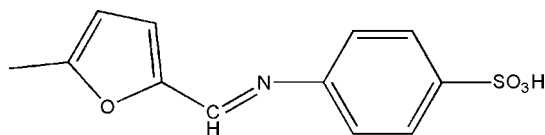
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.045; wR factor = 0.128; data-to-parameter ratio = 13.1.

The title compound, $\text{C}_{12}\text{H}_{11}\text{NO}_4\text{S}$, is a Schiff base derived from the condensation reaction of equimolar quantities of sulfamide and furfural. The molecule has a *trans* configuration with respect to the imine $\text{C}=\text{N}$ double bond. The N atom is involved in an intermolecular $\text{O}-\text{H}\cdots\text{N}$ hydrogen bond.

Related literature

For related literature, see: Abd El Rehim *et al.* (2001); Hariharan & Urbach (1969); Koning & Cantilena (1994); Tarafder *et al.* (2002).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{11}\text{NO}_4\text{S}$
 $M_r = 265.28$
Monoclinic, $P2_1/c$
 $a = 13.9761$ (11) Å

$b = 11.9820$ (15) Å
 $c = 7.3266$ (10) Å
 $\beta = 95.8010$ (10)°
 $V = 1220.6$ (2) Å³

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.27$ mm⁻¹

$T = 298$ (2) K
 $0.23 \times 0.20 \times 0.15$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.940$, $T_{\max} = 0.961$
6042 measured reflections
2156 independent reflections
1581 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.128$
 $S = 1.05$
2156 reflections
165 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.43$ e Å⁻³
 $\Delta\rho_{\min} = -0.44$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{O1}-\text{H1}\cdots\text{N1}^i$	0.82	2.21	3.025 (3)	176

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

Data collection: SMART (Bruker, 2000); cell refinement: SAINT (Bruker, 2000); 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 (Sheldrick, 2008).

The author acknowledges a research grant (No. 08JZ07) from the Phytochemistry Key Laboratory of Shaanxi province.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BX2168).

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supplementary materials

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(*E*)-4-[(5-Methyl-2-furyl)methyleneamino]benzenesulfonic acid

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Comment

In the past decade, Schiff base compounds have been of great interest. These compounds play an important role in the development of coordination chemistry. Some of the complexes derived from Schiff bases have been found to be complexes with pharmacological and antitumor properties (Abd El Rehim *et al.*, 2001; Koning & Cantilena, 1994; Tarafder *et al.*, 2002). As an extension of the work on the structural characterization of Schiff base compounds, the crystal structure of the title compound, (I), is reported here.

The title compound (I) is a sulfamide derivative. The molecular structure is shown in Fig.1. All bond lengths and bond angles are in the normal ranges and comparable to those observed in a similar sulfamide Schiff base (Hariharan & Urbach 1969). The dihedral angle between the benzene ring and the furfural system is 31.9 (6)°. The torsion angles of N1—C4—C5—C6 and N1—C7—C8—C9 are 178.8 (2) ° and -174.2 (3) °, respectively. The molecular structure adopts a *trans* configuration about the C7 = N1 bond. Table 1 shows hydrogen-bond geometry and a packing diagram is shown in Fig.2.

Experimental

Sulfamide (0.1 mmol, 17.2 mg) and furfural(0.1 mmol, 9.6 mg) were dissolved in methanol (10 ml). The mixture was stirred for 30 min at room temperature to give a clear brown solution. After allowing the resulting solution to stand in air for 7 d, brown flake-shaped crystals of (I) were formed on slow evaporation of the solvent. The crystals were collected, washed with methanol and dried in a vacuum desiccator using anhydrous CaCl₂ (yield 54%). Analysis found: C 54.28%, H 4.15% calculated for C₁₂H₁₁NO₄S: C 54.34%, H 4.15%.

Refinement

All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms with C—H distances in the range 0.93–0.96 Å and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}$ or $1.5U_{\text{eq}}(\text{C/O})$

Figures

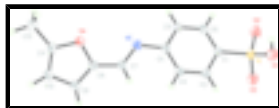


Fig. 1. The structure of the title compound in 30% probability ellipsoids. H atoms are shown as spheres of arbitrary radii.

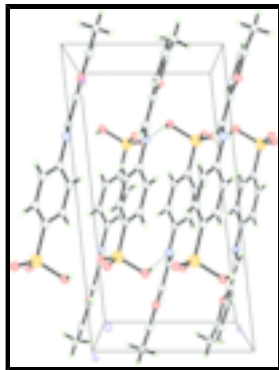


Fig. 2. The molecular packing of (I) viewed along the *b* axis. The dotted lines represent hydrogen bonds. [Symmetry code: (A) $-x + 1, y, -z + 1$]

(E)-4-[(5-Methyl-2-furyl)methyleneamino]benzenesulfonic acid

Crystal data

$C_{12}H_{11}NO_4S$

$M_r = 265.28$

Monoclinic, $P2_1/c$

$a = 13.9761$ (11) Å

$b = 11.9820$ (15) Å

$c = 7.3266$ (10) Å

$\beta = 95.8010$ (10)°

$V = 1220.6$ (2) Å³

$Z = 4$

$F_{000} = 552$

$D_x = 1.444$ Mg m⁻³

Mo $K\alpha$ radiation

$\lambda = 0.71073$ Å

Cell parameters from 1942 reflections

$\theta = 2.9$ – 26.0 °

$\mu = 0.27$ mm⁻¹

$T = 298$ (2) K

Flake, brown

$0.23 \times 0.20 \times 0.15$ mm

Data collection

Bruker SMART CCD area-detector diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

$T = 298$ (2) K

φ and ω scans

Absorption correction: multi-scan (SADABS; Sheldrick, 1996)

$T_{\min} = 0.940$, $T_{\max} = 0.961$

6042 measured reflections

2156 independent reflections

1581 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.028$

$\theta_{\text{max}} = 25.0$ °

$\theta_{\text{min}} = 1.5$ °

$h = -14 \rightarrow 16$

$k = -14 \rightarrow 9$

$l = -8 \rightarrow 8$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.045$

$wR(F^2) = 0.128$

Secondary atom site location: difference Fourier map

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0528P)^2 + 1.108P]$$

where $P = (F_o^2 + 2F_c^2)/3$

$S = 1.05$	$(\Delta/\sigma)_{\max} < 0.001$
2156 reflections	$\Delta\rho_{\max} = 0.44 \text{ e } \text{\AA}^{-3}$
165 parameters	$\Delta\rho_{\min} = -0.44 \text{ e } \text{\AA}^{-3}$
Primary atom site location: structure-invariant direct methods	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.

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

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.70555 (16)	0.62949 (19)	0.4602 (3)	0.0366 (6)
O1	0.22901 (17)	0.5894 (2)	0.3632 (3)	0.0621 (7)
H1	0.2437	0.5289	0.4103	0.093*
O2	0.27564 (15)	0.5149 (2)	0.0717 (3)	0.0596 (7)
O3	0.26181 (15)	0.71745 (19)	0.1207 (3)	0.0581 (7)
O4	0.90263 (14)	0.64516 (16)	0.5898 (3)	0.0409 (5)
S1	0.28904 (5)	0.60909 (6)	0.18999 (10)	0.0392 (2)
C1	0.41271 (19)	0.6154 (2)	0.2690 (3)	0.0323 (6)
C2	0.4477 (2)	0.7072 (2)	0.3712 (4)	0.0379 (7)
H2	0.4062	0.7648	0.3957	0.045*
C3	0.5434 (2)	0.7130 (2)	0.4362 (4)	0.0375 (7)
H3	0.5661	0.7736	0.5073	0.045*
C4	0.60639 (19)	0.6285 (2)	0.3960 (3)	0.0321 (6)
C5	0.5707 (2)	0.5367 (2)	0.2959 (4)	0.0389 (7)
H5	0.6120	0.4790	0.2711	0.047*
C6	0.4741 (2)	0.5301 (2)	0.2326 (4)	0.0389 (7)
H6	0.4508	0.4683	0.1656	0.047*
C7	0.7498 (2)	0.7230 (2)	0.4686 (4)	0.0381 (7)
H7	0.7155	0.7862	0.4270	0.046*
C8	0.8478 (2)	0.7368 (2)	0.5371 (4)	0.0383 (7)
C9	0.9011 (2)	0.8304 (3)	0.5670 (4)	0.0472 (8)
H9	0.8809	0.9033	0.5424	0.057*
C10	0.9932 (2)	0.7963 (3)	0.6424 (4)	0.0476 (8)
H10	1.0452	0.8426	0.6781	0.057*
C11	0.9918 (2)	0.6844 (3)	0.6528 (4)	0.0425 (7)
C12	1.0641 (2)	0.5993 (3)	0.7175 (5)	0.0583 (9)
H12A	1.1232	0.6355	0.7616	0.087*

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H12B	1.0750	0.5505	0.6179	0.087*
H12C	1.0409	0.5568	0.8149	0.087*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.0333 (13)	0.0386 (14)	0.0368 (13)	-0.0029 (10)	-0.0015 (10)	0.0024 (10)
O1	0.0497 (14)	0.0632 (16)	0.0740 (17)	-0.0002 (12)	0.0089 (13)	0.0110 (13)
O2	0.0425 (13)	0.0767 (17)	0.0577 (14)	-0.0077 (11)	-0.0044 (11)	-0.0265 (12)
O3	0.0416 (12)	0.0638 (15)	0.0671 (16)	0.0026 (11)	-0.0034 (11)	0.0290 (12)
O4	0.0364 (11)	0.0362 (11)	0.0490 (12)	-0.0018 (8)	-0.0017 (9)	0.0015 (9)
S1	0.0308 (4)	0.0463 (5)	0.0393 (4)	-0.0031 (3)	-0.0026 (3)	0.0019 (3)
C1	0.0306 (14)	0.0370 (15)	0.0288 (14)	-0.0035 (12)	0.0009 (11)	0.0023 (12)
C2	0.0352 (15)	0.0367 (16)	0.0418 (16)	0.0035 (12)	0.0041 (13)	-0.0049 (13)
C3	0.0384 (16)	0.0369 (16)	0.0362 (15)	-0.0046 (12)	-0.0003 (13)	-0.0077 (12)
C4	0.0321 (14)	0.0342 (15)	0.0293 (14)	-0.0023 (11)	-0.0008 (11)	0.0039 (11)
C5	0.0334 (15)	0.0353 (16)	0.0476 (17)	0.0026 (12)	0.0017 (13)	-0.0045 (13)
C6	0.0380 (16)	0.0345 (16)	0.0433 (17)	-0.0046 (12)	-0.0003 (13)	-0.0075 (13)
C7	0.0354 (15)	0.0371 (16)	0.0406 (16)	-0.0012 (13)	-0.0013 (13)	0.0033 (13)
C8	0.0375 (16)	0.0371 (16)	0.0394 (16)	-0.0013 (12)	-0.0012 (13)	0.0025 (13)
C9	0.0441 (18)	0.0349 (17)	0.060 (2)	-0.0035 (14)	-0.0054 (15)	0.0040 (15)
C10	0.0363 (17)	0.0469 (19)	0.057 (2)	-0.0104 (14)	-0.0080 (15)	-0.0009 (15)
C11	0.0327 (16)	0.0493 (19)	0.0443 (17)	-0.0026 (13)	-0.0020 (13)	0.0005 (14)
C12	0.0444 (19)	0.057 (2)	0.072 (2)	0.0109 (16)	-0.0017 (17)	0.0028 (18)

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

N1—C7	1.278 (3)	C4—C5	1.386 (4)
N1—C4	1.418 (3)	C5—C6	1.385 (4)
O1—S1	1.608 (2)	C5—H5	0.9300
O1—H1	0.8200	C6—H6	0.9300
O2—S1	1.424 (2)	C7—C8	1.420 (4)
O3—S1	1.431 (2)	C7—H7	0.9300
O4—C11	1.368 (3)	C8—C9	1.352 (4)
O4—C8	1.372 (3)	C9—C10	1.409 (4)
S1—C1	1.768 (3)	C9—H9	0.9300
C1—C6	1.377 (4)	C10—C11	1.344 (4)
C1—C2	1.392 (4)	C10—H10	0.9300
C2—C3	1.375 (4)	C11—C12	1.478 (4)
C2—H2	0.9300	C12—H12A	0.9600
C3—C4	1.394 (4)	C12—H12B	0.9600
C3—H3	0.9300	C12—H12C	0.9600
C7—N1—C4	118.4 (2)	C1—C6—C5	119.9 (3)
S1—O1—H1	109.5	C1—C6—H6	120.0
C11—O4—C8	106.5 (2)	C5—C6—H6	120.0
O2—S1—O3	119.29 (15)	N1—C7—C8	124.3 (3)
O2—S1—O1	108.48 (14)	N1—C7—H7	117.8
O3—S1—O1	105.76 (13)	C8—C7—H7	117.8

O2—S1—C1	107.28 (13)	C9—C8—O4	109.6 (2)
O3—S1—C1	107.07 (13)	C9—C8—C7	130.5 (3)
O1—S1—C1	108.60 (13)	O4—C8—C7	119.9 (2)
C6—C1—C2	119.9 (3)	C8—C9—C10	106.9 (3)
C6—C1—S1	120.9 (2)	C8—C9—H9	126.6
C2—C1—S1	119.2 (2)	C10—C9—H9	126.6
C3—C2—C1	120.2 (3)	C11—C10—C9	107.1 (3)
C3—C2—H2	119.9	C11—C10—H10	126.5
C1—C2—H2	119.9	C9—C10—H10	126.5
C2—C3—C4	120.2 (3)	C10—C11—O4	110.0 (3)
C2—C3—H3	119.9	C10—C11—C12	133.9 (3)
C4—C3—H3	119.9	O4—C11—C12	116.1 (3)
C5—C4—C3	119.2 (3)	C11—C12—H12A	109.5
C5—C4—N1	118.0 (2)	C11—C12—H12B	109.5
C3—C4—N1	122.8 (2)	H12A—C12—H12B	109.5
C6—C5—C4	120.6 (3)	C11—C12—H12C	109.5
C6—C5—H5	119.7	H12A—C12—H12C	109.5
C4—C5—H5	119.7	H12B—C12—H12C	109.5
O2—S1—C1—C6	-8.0 (3)	C2—C1—C6—C5	-0.8 (4)
O3—S1—C1—C6	-137.2 (2)	S1—C1—C6—C5	-179.9 (2)
O1—S1—C1—C6	109.0 (2)	C4—C5—C6—C1	0.0 (4)
O2—S1—C1—C2	172.8 (2)	C4—N1—C7—C8	177.3 (3)
O3—S1—C1—C2	43.7 (3)	C11—O4—C8—C9	-0.2 (3)
O1—S1—C1—C2	-70.1 (2)	C11—O4—C8—C7	-178.6 (3)
C6—C1—C2—C3	-0.1 (4)	N1—C7—C8—C9	-174.2 (3)
S1—C1—C2—C3	179.1 (2)	N1—C7—C8—O4	3.8 (4)
C1—C2—C3—C4	1.7 (4)	O4—C8—C9—C10	-0.3 (4)
C2—C3—C4—C5	-2.4 (4)	C7—C8—C9—C10	177.9 (3)
C2—C3—C4—N1	-179.6 (2)	C8—C9—C10—C11	0.6 (4)
C7—N1—C4—C5	146.0 (3)	C9—C10—C11—O4	-0.8 (4)
C7—N1—C4—C3	-36.9 (4)	C9—C10—C11—C12	179.8 (4)
C3—C4—C5—C6	1.6 (4)	C8—O4—C11—C10	0.7 (3)
N1—C4—C5—C6	178.8 (2)	C8—O4—C11—C12	-179.8 (3)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
O1—H1 \cdots N1 ⁱ	0.82	2.21	3.025 (3)	176

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

Fig. 1

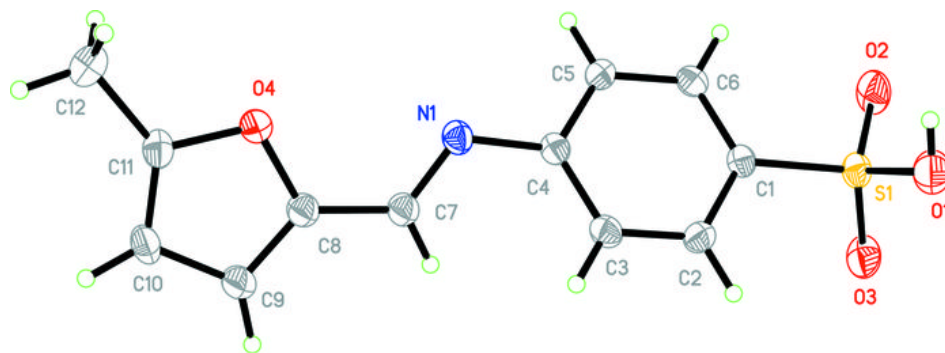


Fig. 2

