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#### **Key indicators**

Single-crystal X-ray study T = 292 K Mean  $\sigma$ (C–C) = 0.004 Å R factor = 0.048 wR factor = 0.122 Data-to-parameter ratio = 18.0

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

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# 2-Chloro-4-fluorophenyl methanesulfonate

In the crystal structure of the title compound, C<sub>7</sub>H<sub>6</sub>ClFO<sub>3</sub>S, molecules form a supramolecular structure via intermolecular C-H···O hydrogen bonds and  $\pi$ - $\pi$  stacking interactions.

#### Comment

The title compound, (I), is an unexpected product of the reaction of 7-chloro-5-methylsulfanyl-1,2,4-triazolo[1,5-c]pyrimidin-2-yl methanesulfonate with 2-chloro-4-fluorophenol.



The molecular structure of (I) is shown in Fig. 1. Selected bond lengths and angels are shown in Table 1. In the crystal structure, molecules are linked by  $C-H \cdots O$  hydrogen bonds (Table 2) to form a chain along the c axis (Fig. 2). The packing is further stabilized by  $\pi - \pi$  stacking interactions between the benzene rings of neighboring molecular chains; the distance between centroids  $[Cg \cdots Cg(1-x, 1-y, 1-z)]$  is 3.684 (2) Å and the interplanar spacing is 3.477(2) Å.

# **Experimental**

A mixture of 2-chloro-4-fluorophenol (4 mmol) and sodium hydride (4 mmol) in anhydrous toluene (60 ml) was stirred at 373 K for 2 h. 7-Chloro-5-methylsulfanyl-1,2,4-triazolo[1,5-c]pyrimidin-2-yl methanesulfonate (1 mmol) was then added and the resulting reaction mixture was refluxed for about 2 h. After filtration, the solvent was removed under reduced pressure and the residue was purified by column chromatography on silical gel (eluent: petroleum-acetone 4:1 v/v) to afford the compound (I) (yield 45%, m.p. 362 K). <sup>1</sup>H NMR (CDCl<sub>3</sub>, 400 MHz): § 7.04–7.44 (m, 3H, Ar-H), 3.25 (s, 3H, CH<sub>3</sub>) Crystals suitable for X-ray analysis were grown from a diethyl ethyl ether solution at 277 K.

Crystal data C7H6ClFO3S  $D_r = 1.651 \text{ Mg m}^{-3}$ Mo  $K\alpha$  radiation  $M_r = 224.63$ Monoclinic,  $P2_1/c$ a = 11.2684 (13) Åreflections b = 7.9014 (9) Å  $\theta = 3.3 - 21.9^{\circ}$  $\mu = 0.64 \text{ mm}^{-1}$ c = 11.1345 (12) Å $\beta = 114.284 \ (2)^{\circ}$ T = 292 (2) K Plate, colorless  $V = 903.65 (18) \text{ Å}^3$ Z = 4

# organic papers

### Data collection

Bruker SMART 4K CCD area-	2138 independent reflections
detector diffractometer	1467 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans	$R_{\rm int} = 0.038$
Absorption correction: multi-scan	$\theta_{\rm max} = 28.0^{\circ}$
(SADABS; Sheldrick, 1996)	$h = -14 \rightarrow 14$
$T_{\min} = 0.832, \ T_{\max} = 0.963$	$k = -10 \rightarrow 10$
7603 measured reflections	$l = -13 \rightarrow 14$

#### Refinement

Refinement on  $F^2$  $R[F^2 > 2\sigma(F^2)] = 0.048$ wR(F<sup>2</sup>) = 0.122 S = 1.042138 reflections 119 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

C1-Cl1	1.720 (3)	O1-S1	1.5996 (18)
C3-F1	1.353 (3)	C7-S1	1.736 (3)
C6-O1	1.407 (3)		
C6-O1-S1	120.05 (15)		
C6-O1-S1-C7	88.1 (2)		

+ 0.1158P] where  $P = (F_0^2 + 2F_c^2)/3$ 

 $(\Delta/\sigma)_{\rm max} < 0.001$  $\Delta \rho_{\rm max} = 0.28 \text{ e } \text{\AA}^{-3}$ 

 $\Delta \rho_{\rm min} = -0.27 \text{ e} \text{ Å}^{-3}$ 

 $w = 1/[\sigma^2(F_0^2) + (0.0553P)^2]$ 

# Table 2

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdot \cdot \cdot A$
$C7-H7A\cdots O4^{i}$	0.96	2.42	3.318 (3)	155
Symmetry code: (i) x	$y - y + \frac{1}{2}, z - \frac{1}{2}.$			

H atoms were placed at calculated positions and treated as riding atoms, with C-H = 0.93–0.96 Å and  $U_{iso}(H) = 1.2U_{eq}(\text{aromatic C})$  or  $1.5U_{eq}$ (methyl C).

Data collection: SMART (Bruker, 1997); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2001); software used to prepare material for publication: SHELXTL.

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Figure 1

A view of (I), showing the atom-labellng scheme. Displacement ellipsoids are drawn at the 50% probability level.



### Figure 2

Packing of (I), viewed down the b axis, showing  $\pi - \pi$  stacking interactions (dotted lines) and C-H···O hydrogen bonds (dashed lines). The symmetry code is as in Table 2.

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