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Key indicators

Single-crystal X-ray study T = 150 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.034 wR factor = 0.086 Data-to-parameter ratio = 17.8

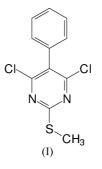
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

4,6-Dichloro-2-methylthio-5-phenylpyrimidine

The structure of the title compound, $C_{11}H_8Cl_2N_2S$, comprises two unique molecules with the dihedral angles between the two linked rings for each molecule being 77.1 (1) and 78.4 (1)°. Only one $C-H\cdots X$ short contact exists from an aromatic C-H group to a Cl atom.

Comment

The title compound, (I), is an analogue of 4,6-dichloro-2methylthiopyrimidine (dcmtp; Lynch & McClenaghan, 2000). In dcmtp, the Cl atoms are displaced by nucleophiles first, before the methylthio group, although the second Cl atom is much less reactive than the first. In terms of substituent effects, the presence of the phenyl ring in (I) should increase the activity of the Cl atoms, but as found for 4,6-dichloro-2-amino-5-phenylpyrimidine, nucleophilic displacement of the first Cl atom takes a reaction time of hours, whereas that of the second takes days. This rate difference may be due to steric effects, thus we have instigated a series of studies to examine the structural changes that the phenyl ring imposes on the nucleophilic substituents of several different analogues of (I), including the 2-amino series. Here we report the single-crystal structure of (I) as our point of reference for further derivatized compounds in the 2-methylthio series.



Experimental

The title compound, (I), was prepared by Spa Contract Synthesis. Crystals of (I) formed when the reaction solution $(POCl_3)$ was poured into excess water.

Crystal data

C₁₁H₈Cl₂N₂S $M_r = 271.15$ Orthorhombic, $P2_12_12_1$ a = 7.467 (2) Å b = 16.217 (3) Å c = 19.318 (4) Å V = 2339.1 (8) Å³ Z = 8 $D_x = 1.540$ Mg m⁻³ Mo $K\alpha$ radiation Cell parameters from 9258 reflections $\theta = 1.0-27.5^{\circ}$ $\mu = 0.70 \text{ mm}^{-1}$ T = 150 (2) KPrism, colourless $0.28 \times 0.24 \times 0.20 \text{ mm}$

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Data collection

Enraf–Nonius KappaCCD area- detector diffractometer φ and ω scans Absorption correction: multi-scan (<i>SORTAV</i> ; Blessing, 1995) $T_{\min} = 0.827, T_{\max} = 0.872$ 11 464 measured reflections	5186 independent reflections 4655 reflections with $I > 2\sigma(I)$ $R_{int} = 0.045$ $\theta_{max} = 27.5^{\circ}$ $h = -9 \rightarrow 9$ $k = -19 \rightarrow 20$ $l = -24 \rightarrow 24$
Refinement	
Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.034$	$w = 1/[\sigma^2(F_o^2) + (0.0400P)^2 + 0.3307P]$
$wR(F^2) = 0.087$ S = 1.06 5186 reflections	where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\text{max}} = 0.010$ $\Delta\rho_{\text{max}} = 0.23 \text{ e } \text{\AA}^{-3}$
291 parameters	$\Delta \rho_{\rm min} = -0.31 \text{ e} \text{ Å}^{-3}$

Table 1

Selected geometric parameters (°).

H-atom parameters constrained

N1A-C2A-S21A-C22A	3.7 (2)	N1B-C2B-S21B-C22B	3.4 (2)
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Absolute structure: Flack (1983)

Flack parameter = -0.04(5)

Table 2

Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$C22B - H22B \cdot \cdot \cdot Cl4B^{i}$	0.98	2.76	3.452 (3)	128
Symmetry code: (i) $\frac{3}{2} - x$, 1	$1 - y, \frac{1}{2} + z.$			

All H atoms were included in the refinement at calculated positions as riding models with C–H distances set to 0.95 (Ar-H) and 0.98 Å (CH₃). The number of Friedel pairs is 2167.

Data collection: *DENZO* (Otwinowski & Minor, 1997) and *COLLECT* (Hooft, 1998); cell refinement: *DENZO* and *COLLECT*; data reduction: *DENZO* and *COLLECT*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics:

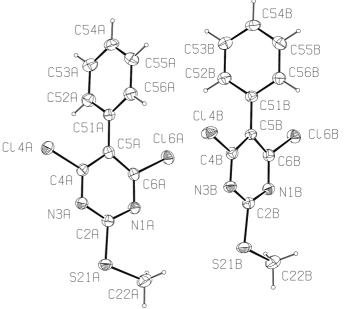


Figure 1

The molecular configuration and atom-numbering scheme for (I), showing 50% probability ellipsoids.

*PLATON*97 (Spek, 1997); software used to prepare material for publication: *SHELXL*97.

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