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## Structure Reports

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## 6-Chloro-4-(dimethylaminomethyleneamino)-2-(methylsulfanyl)pyrimidine

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## Key indicators

Single-crystal X-ray study
$T=120 \mathrm{~K}$
Mean $\sigma(\mathrm{C}-\mathrm{C})=0.003 \AA$
$R$ factor $=0.040$
$w R$ factor $=0.106$
Data-to-parameter ratio $=18.3$

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

[^0]The molecules in the title compound, $\mathrm{C}_{8} \mathrm{H}_{11} \mathrm{ClN}_{4} \mathrm{~S}$, are linked in pairs by a $\pi-\pi$ stacking interaction. There are, however, no other direction-specific interactions.

## Comment

In our search for good candidates for intermediates in the synthesis of new pyrimidine fused ring systems, we have prepared the title compound, (I), (Fig. 1), a formyl derivative of 4-amino-6-chloro-2-(methylsulfanyl)pyrimidine, using the Vilsmeier formylation reaction (Vilsmeier \& Haack, 1927).

(I)

The bond lengths and angles show no unusual features. The essentially planar group consisting of atoms N4, C41, N42, C43 and C44 forms a dihedral angle of 31.49 (8) ${ }^{\circ}$ with that of the planar pyrimidine ring. The leading torsion angles are given in Table 1. The molecules are linked into pairs by a $\pi-\pi$ stacking


Figure 1
A view of (I) with our numbering scheme. Displacement ellipsoids are drawn at the $30 \%$ probability level.

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interaction (Fig. 2). The molecules at $(x, y, z)$ and ( $1-x, 1-y$, $1-z$ ) are parallel, with an interplanar spacing of 3.4661 (2) A. The ring-centroid separation is 3.359 (2) $\AA$ corresponding to a ring offset of $0.857 \AA$.

## Experimental

The Vilsmeier reagent was prepared in an ice-bath by adding phosphorus oxychloride $(1.8 \mathrm{mmol})$ to $N, N$-dimethylformamide ( 38 mmol ) and stirring for 15 min . 4-Amino-6-chloro-2-(methylsulfanyl)pyrimidine ( $0.2 \mathrm{~g}, 1.14 \mathrm{mmol}$ ) was then added and the reaction temperature raised to $323-333 \mathrm{~K}$, and the mixture stirred for 2 h . The reaction mixture was then poured on to crushed ice and neutralized with NaOH ( $10 \%$ in water) until the pH was raised to $8-$ 9. The resulting white solid was filtered off and recrystallized from DMSO- $d_{6}$ producing white crystalline blocks suitable for singlecrystal X-ray diffraction (yield $60 \%$; m.p. 374-376 K). MS ( 70 eV ): 232/230 (38:100, $\left.M+2 / M^{+}\right)$, 217/215 (17/18, $\left.\left[(M+2 / M)-\mathrm{CH}_{3}\right]^{+}\right), 186 /$ $184\left(17 / 18,\left[(M+2 / M)-\mathrm{SCH}_{2}\right]^{+}\right), 149\left(31,\left[M-\mathrm{SCH}_{3}-\mathrm{Cl}\right]^{+}\right), 71$ (4, $\left.\left[\mathrm{N}=\mathrm{CH}-\mathrm{N}\left(\mathrm{CH}_{3}\right)_{2}\right]^{+}\right)$.

## Crystal data

$\mathrm{C}_{8} \mathrm{H}_{11} \mathrm{ClN}_{4} \mathrm{~S}$
$M_{r}=230.72$
Triclinic, $P \overline{1}$
$a=7.4817(2) \AA$
$b=8.5739$ (2) $\AA$
$c=9.818$ (3) A
$\alpha=111.973$ (2) ${ }^{\circ}$
$\beta=91.661$ (2) ${ }^{\circ}$
$\gamma=114.566(2)^{\circ}$

## Data collection

Nonius KappaCCD diffractometer $\varphi$ and $\omega$ scans
Absorption correction: multi-scan
(SADABS; Sheldrick, 2003)
$T_{\text {min }}=0.856, T_{\text {max }}=0.901$

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.040$
$w R\left(F^{2}\right)=0.106$
$S=1.13$
2378 reflections
130 parameters
H -atom parameters constrained

$$
V=518.31(15) \AA^{3}
$$

$Z=2$
$D_{x}=1.478 \mathrm{Mg} \mathrm{m}^{-3}$
Mo $K \alpha$ radiation
$\mu=0.54 \mathrm{~mm}^{-1}$
$T=120$ (2) K
Block, colourless
$0.30 \times 0.30 \times 0.20 \mathrm{~mm}$

12192 measured reflections 2378 independent reflections 2015 reflections with $I>2 \sigma(I)$ $R_{\text {int }}=0.032$
$\theta_{\text {max }}=27.5^{\circ}$

$$
\begin{aligned}
& w=1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0456 P)^{2}\right. \\
& \quad+0.5109 P] \\
& \text { where } P=\left(F_{\mathrm{o}}{ }^{2}+2 F_{\mathrm{c}}{ }^{2}\right) / 3 \\
& (\Delta / \sigma)_{\max }<0.001 \\
& \Delta \rho_{\max }=0.43 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.36 \mathrm{e}^{-3}
\end{aligned}
$$

Table 1
Selected torsion angles $\left({ }^{\circ}\right)$.

| N3-C2-S2-C21 | $0.17(18)$ | N4-C41-N41-C43 | $-3.4(3)$ |
| :--- | :---: | :--- | ---: |
| N1-C2-S2-C21 | $-179.75(14)$ | N4-C41-N41-C44 | $175.22(18)$ |
| N3-C4-N4-C41 | $-25.4(3)$ | C2-N1-C6-Cl6 | $-177.26(13)$ |
| C5-C4-N4-C41 | $156.38(18)$ | C4-C5-C6-Cl6 | $175.75(14)$ |
| C4-N4-C41-N41 | $174.23(17)$ |  |  |

H atoms were treated as riding atoms, with aromatic $\mathrm{C}-\mathrm{H}=$ $0.95 \AA$ and $U_{\text {iso }}(\mathrm{H})=1.2 U_{\text {eq }}(\mathrm{C})$, and $\mathrm{C}-\mathrm{H}=0.98 \AA$ and $U_{\text {iso }}(\mathrm{H})=$ $1.5 U_{\text {eq }}(\mathrm{C})$. The positions of all methyl H atoms were checked in a difference map.

Data collection: COLLECT (Bruker-Nonius, 2004); cell refinement: DIRAX/LSQ (Duisenberg et al., 2000); data reduction:


Figure 2
A view of the $\pi-\pi$ stacking viewed perpendicular to the plane of the pyrimidine ring. Atoms labelled with an asterisk (*) are in the molecule at $(1-x, 1-y, 1-z)$. For the sake of clarity, all H atoms have been omitted.

EVALCCD (Duisenberg et al., 2003); program(s) used to solve structure: SIR2004 (Burla et al., 2005); program(s) used to refine structure: OSCAIL (McArdle, 2003) and SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97 and WORDPERFECT macro PRPKAPPA (Ferguson, 1999).

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