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Li Tian^a* and Da-Xiang Wang^b

^aCollege of Chemistry and Life Sciences, Tianjin Normal University, Tianjin 300074, People's Republic of China, and ^bChina Automotive Technology and Research Centre, PO Box 59, Tianjin 300162, People's Republic of China

Correspondence e-mail: tian_lili@eyou.com

Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.005 Å R factor = 0.039 wR factor = 0.121 Data-to-parameter ratio = 12.4

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

6-Cyano-3,3-dimethyl-5-methylsulfanyl-1,3-dihydro-2-oxo-1-thia-3b,4,8-triazaindacene 1,1-dioxide

The title compound, $C_{11}H_{10}N_4O_3S_2$, has normal bond lengths and angles. The molecule includes a pyrazole ring, a pyrimidine ring and a sulfone ring. The crystal packing is mainly stabilized by van der Waals interactions. Received 2 March 2005 Accepted 4 April 2005 Online 9 April 2005

Comment

Vilsmeier reagents, HCON R_1R_2 /POCl₃, are extensively used in the synthesis of aldehyde derivatives and formamidines (Meth-Cohn, 1991). Recently, it has been found that application of the Vilsmeier reaction with acetylphosphonates leads stereospecifically to (*Z*)- β -phosphonyl- β -chlorovinylaldehydes (Qian *et al.*, 2000). Chloroformylation of β -carbonyl sulfone (Ingate *et al.*, 1997) with the Vilsmeier reagent DMF/ POCl₃ afforded the cyclic β -chlorovinylaldehyde 4-chloro-5,5dimethyl-3-formyl-1,2-oxathiolene 2,2-dioxide, (1). This compound is a very useful intermediate for the synthesis of heterocyclic compounds, and reaction with 5-amino-4-cyano-3-methylsulfanyl pyrazole, (2), gives the title heterocyclic compound, (3). We report here the crystal structure of (3).



The molecular structure of (3) is shown in Fig. 1. All bond lengths and angles are normal (Table 1). The crystal packing



© 2005 International Union of Crystallography Printed in Great Britain – all rights reserved A view of the title compound, showing displacement ellipsoids at the 40% probability level.





(Fig. 2) is stabilized by van der Waals interactions. As depicted in Fig. 1, there is a six-membered pyrimidine ring, N1/C4/C5/ C6/N2/C7, which is formed by the amino group of (2) bonded with atom C4 of (1) and the imino group of (2) bonded with the formyl group of (1), rather than a combination of the amino group of (2) combined with the formyl group of (1), as has been previously observed by Tian & Liu (2004). The short bond lengths of C5-C6, C7-C8 and C4-N1 (Table 1) are indicative of significant double-bond character.

Experimental

A solution of 5-amino-4-cyano-3-methylsulfanyl pyrazole, (2) (1 mmol, 0.16 g), in water (2 ml) was added dropwise to a solution of compound (1) (1 mmol, 0.21 g) in dichloromethane (5 ml) at 273-283 K. K₂CO₃ (1 mmol, 0.14 g) in water (2 ml) was then added dropwise at this temperature. The reaction mixture was kept at room temperature for 2-3 h. The aqueous layer was extracted with dichloromethane and the combined organic layers were washed with saturated brine, dried, filtered and concentrated. The residue was separated by silica gel to afford the title compound. Recrystallization from a solution in ethyl acetate-cyclohexane (1:3) yielded colourless single crystals suitable for X-ray diffraction studies. ¹H NMR (300 MHz, CDCl₃, δ, p.p.m.): 7.05 (s, 1H, H9), 2.75 (s, 3H, -SCH₃), 1.86 [s, 6H, (CH₃)₂]. Analysis, calculated for C₁₁H₁₀N₄S₂O₃: C 42.57, H 3.25, N 18.05%; found: C 42.40, H 3.39, N 17.98%.

Crystal data

$C_{11}H_{10}N_4O_3S_2$	Z = 2
$M_r = 310.35$	$D_x = 1.564 \text{ Mg m}^{-3}$
Triclinic, $P\overline{1}$	Mo $K\alpha$ radiation
a = 7.150 (2) Å	Cell parameters from 1868
b = 8.573 (3) Å	reflections
c = 11.171 (3) Å	$\theta = 2.4-27.9^{\circ}$
$\alpha = 98.316 \ (4)^{\circ}$	$\mu = 0.42 \text{ mm}^{-1}$
$\beta = 98.953 \ (4)^{\circ}$	T = 293 (2) K
$\gamma = 98.420 \ (4)^{\circ}$	Prism, colourless
V = 659.2 (3) Å ³	$0.38 \times 0.16 \times 0.10 \ \text{mm}$

Data collection

Bruker APEX II CCD area detector diffractometer φ and ω scans Absorption correction: multi-scan (<i>SADABS</i> ; Sheldrick, 1996) $T_{min} = 0.784, T_{max} = 0.960$ 3565 measured reflections	2279 independent reflections 1954 reflections with $I > 2\sigma(I)$ $R_{int} = 0.015$ $\theta_{max} = 25.0^{\circ}$ $h = -8 \rightarrow 8$ $k = -10 \rightarrow 9$ $l = -13 \rightarrow 12$
Refinement	
Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.039$ $wR(F^2) = 0.121$ S = 1.08 2279 reflections 184 parameters H-atom parameters constrained	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0487P)^{2} + 0.9464P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.29 \text{ e } \text{ Å}^{-3}$ $\Delta\rho_{min} = -0.33 \text{ e } \text{ Å}^{-3}$

Table 1 Selected geometric parameters (Å, °).

N1-C4	1.315 (4)	N3-C10	1.331 (4)
N1-C7	1.353 (4)	C1-C3	1.513 (5)
N2-C6	1.338 (4)	C4-C5	1.400 (4)
N2-N3	1.374 (3)	C5-C6	1.365 (4)
N2-C7	1.393 (4)	C7-C8	1.377 (4)
03 - 81 - C5	93.66 (13)	C6 - C5 - C4	120.2 (3)
C4-N1-C7	114.5 (2)	C6-C5-S1	130.8 (2)
C6-N2-N3	124.6 (2)	C4-C5-S1	109.0(2)
C6-N2-C7	123.0 (2)	N2-C6-C5	115.2 (3)
N3-N2-C7	112.4 (2)	N1-C7-C8	132.4 (3)
C10-N3-N2	103.9 (2)	N1-C7-N2	122.1 (2)
O3-C3-C4	102.8 (2)	C8-C7-N2	105.5 (2)
N1-C4-C5	124.9 (3)	C7-C8-C10	105.6 (3)
N1-C4-C3	122.0 (3)	N3-C10-C8	112.6 (3)
C5-C4-C3	113.1 (3)		

All H atoms were placed in calculated positions, with C-H = 0.93or 0.96 Å, and included in the final cycles of refinement using a riding model, with $U_{iso}(H) = 1.2U_{eq}(C)$ [1.5 U_{eq} for methyl].

Data collection: APEXII (Bruker, 1998); cell refinement: APEXII; data reduction: SAINT (Bruker, 1999); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 1999); software used to prepare material for publication: SHELXTL.

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