

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

3-[1-(4-Chlorophenyl)ethyl]-1,3-thiazinane-2-thione

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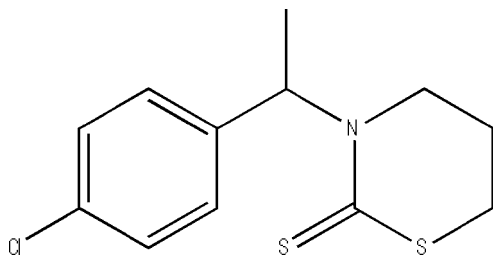
Received 6 January 2011; accepted 13 January 2011

Key indicators: single-crystal X-ray study; $T = 113$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.038; wR factor = 0.089; data-to-parameter ratio = 19.9.

In the title compound, $\text{C}_{12}\text{H}_{14}\text{ClNS}_2$, the thiazole ring adopts an envelope conformation; the basal plane is nearly perpendicular to the benzene ring at a dihedral angle of $85.72(5)^\circ$. Weak intermolecular $\text{C}-\text{H}\cdots\text{S}$ hydrogen bonding is present in the crystal structure.

Related literature

For the biological activity of thiazole compounds, see: Amir *et al.* (2006). For a related structure, see: Cunico *et al.* (2007).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{14}\text{ClNS}_2$

$M_r = 271.81$

Orthorhombic, $Pbca$

$a = 11.260(2)$ Å

$b = 11.888(2)$ Å

$c = 18.978(4)$ Å

$V = 2540.5(9)$ Å³

$Z = 8$

Mo $K\alpha$ radiation

$\mu = 0.60$ mm⁻¹

$T = 113$ K

$0.18 \times 0.14 \times 0.12$ mm

Data collection

Rigaku Saturn diffractometer

Absorption correction: multi-scan

(*CrystalClear*; Rigaku, 2005)

$T_{\min} = 0.900$, $T_{\max} = 0.931$

16988 measured reflections

2932 independent reflections

2605 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.050$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.038$

$wR(F^2) = 0.089$

$S = 1.11$

2925 reflections

147 parameters

H-atom parameters constrained

$\Delta\rho_{\text{max}} = 0.29$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.26$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C9}-\text{H9A}\cdots\text{S2}^i$	0.97	2.85	3.773 (2)	158
$\text{C10}-\text{H10B}\cdots\text{S2}^{ii}$	0.97	2.77	3.701 (2)	160

Symmetry codes: (i) $x + \frac{1}{2}, y, -z + \frac{1}{2}$; (ii) $-x, y + \frac{1}{2}, -z + \frac{1}{2}$.

Data collection: *CrystalClear* (Rigaku, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL*.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5140).

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supplementary materials

Acta Cryst. (2011). E67, o514 [doi:10.1107/S1600536811002078]

3-[1-(4-Chlorophenyl)ethyl]-1,3-thiazinane-2-thione

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Comment

Recently, compounds containing a 1,3-thiazinane group have attracted much interest because the 1,3-thiazinane ring system are well known as its efficient insecticidal activity for a wide variety of crops (Amir *et al.*, 2006). The title compound (I) was synthesized as a new compound with better biological activity. We report here the crystal structure of (I).

In (I) all bond lengths and angles are normal and in a good agreement with those reported previously (Cunico *et al.*, 2007). The thiazole ring is in an envelope conformation with the $-\text{CH}_2-$ group bonded to the S1 atom forming the flap. The 1,3-thiazinane-2-thione ring forms two dihedral angles are $85.99(2)^\circ$ [S1/S2/N1/C7/C9/C11/C12] and $77.68(2)^\circ$ [N1/C9/C10/C11/C12] with the benzene ring respectively. The crystal structure is stabilized by weak intermolecular C–H \cdots S hydrogen bonds.

Experimental

1,3-Thiazinane-2-thione 1.33 g (10.0 mmol) and deacid reagent potassium carbonate 1.38 g (5.0 mmol) were added in a flask equipped with stirrer, the solvent acetonitrile (20 ml) was added and the mixture was stirred for 0.5 h. Then 1-chloro-4-(1-chloroethyl)benzene 1.74 g (10.0 mmol) was added dropwisely within 2 h at 333 K. The mixture was stirred for 8 h at 433 K. Upon cooling at room temperature, then the solid was filtered, the filter-cake was washed twice by acetonitrile. Crystallized from methanol to afford the title compound 2.0 g (74% yield) Single crystals suitable for X-ray measurement were obtained by recrystallization from the mixture of acetone and methanol at room temperature.

Refinement

H atoms were placed in calculated positions, with C–H = 0.93–0.98 Å, and included in the final cycles of refinement using a riding model with $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$ for methyl H atoms and $1.2U_{\text{eq}}(\text{C})$ for the others.

Figures

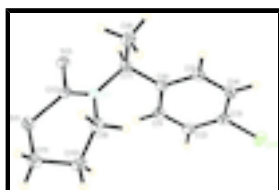


Fig. 1. The molecular structure of (I), with displacement ellipsoids drawn at the 40% probability level.

3-[1-(4-Chlorophenyl)ethyl]-1,3-thiazinane-2-thione

Crystal data

$C_{12}H_{14}ClNS_2$	$F(000) = 1136$
$M_r = 271.81$	$D_x = 1.421 \text{ Mg m}^{-3}$
Orthorhombic, <i>Pbca</i>	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
Hall symbol: -P 2ac 2ab	Cell parameters from 5973 reflections
$a = 11.260 (2) \text{ \AA}$	$\theta = 2.2\text{--}27.5^\circ$
$b = 11.888 (2) \text{ \AA}$	$\mu = 0.60 \text{ mm}^{-1}$
$c = 18.978 (4) \text{ \AA}$	$T = 113 \text{ K}$
$V = 2540.5 (9) \text{ \AA}^3$	Block, colorless
$Z = 8$	$0.18 \times 0.14 \times 0.12 \text{ mm}$

Data collection

Rigaku Saturn diffractometer	2932 independent reflections
Radiation source: rotating anode confocal	2605 reflections with $I > 2\sigma(I)$
ω scans	$R_{\text{int}} = 0.050$
Absorption correction: multi-scan (<i>CrystalClear</i> ; Rigaku, 2005)	$\theta_{\text{max}} = 27.5^\circ$, $\theta_{\text{min}} = 2.2^\circ$
$T_{\text{min}} = 0.900$, $T_{\text{max}} = 0.931$	$h = -14 \rightarrow 14$
16988 measured reflections	$k = -7 \rightarrow 15$
	$l = -24 \rightarrow 24$

Refinement

Refinement on F^2	Secondary atom site location: difference Fourier map
Least-squares matrix: full	Hydrogen site location: inferred from neighbouring sites
$R[F^2 > 2\sigma(F^2)] = 0.038$	H-atom parameters constrained
$wR(F^2) = 0.089$	$w = 1/[\sigma^2(F_o^2) + (0.0383P)^2 + 0.9612P]$
$S = 1.11$	where $P = (F_o^2 + 2F_c^2)/3$
2925 reflections	$(\Delta/\sigma)_{\text{max}} = 0.002$
147 parameters	$\Delta\rho_{\text{max}} = 0.29 \text{ e \AA}^{-3}$
0 restraints	$\Delta\rho_{\text{min}} = -0.26 \text{ e \AA}^{-3}$
Primary atom site location: structure-invariant direct methods	Extinction correction: <i>SHELXTL</i> (Sheldrick, 2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
	Extinction coefficient: 0.0031 (5)

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	$U_{\text{iso}}^*/U_{\text{eq}}$
Cl1	-0.02163 (5)	0.84121 (4)	0.54940 (3)	0.03747 (15)
S1	0.12139 (4)	0.46585 (4)	0.16831 (2)	0.02721 (14)
S2	-0.00573 (4)	0.34132 (4)	0.27142 (2)	0.02514 (13)
N1	0.17516 (12)	0.47749 (11)	0.30690 (7)	0.0190 (3)
C1	0.03561 (15)	0.61986 (15)	0.39541 (9)	0.0223 (4)
H1	0.0095	0.6085	0.3495	0.027*
C2	-0.00917 (15)	0.70925 (15)	0.43370 (10)	0.0244 (4)
H2	-0.0652	0.7573	0.4140	0.029*
C3	0.03060 (16)	0.72615 (14)	0.50171 (10)	0.0247 (4)
C4	0.11393 (16)	0.65588 (15)	0.53172 (9)	0.0263 (4)
H4	0.1405	0.6684	0.5774	0.032*
C5	0.15754 (16)	0.56606 (15)	0.49268 (9)	0.0239 (4)
H5	0.2135	0.5183	0.5127	0.029*
C6	0.11905 (14)	0.54630 (13)	0.42412 (9)	0.0191 (3)
C7	0.15969 (15)	0.44518 (14)	0.38191 (9)	0.0211 (4)
H7	0.0951	0.3900	0.3836	0.025*
C8	0.27095 (17)	0.38623 (17)	0.40901 (10)	0.0317 (4)
H8A	0.2939	0.3284	0.3765	0.047*
H8B	0.2548	0.3533	0.4542	0.047*
H8C	0.3342	0.4399	0.4135	0.047*
C9	0.27378 (15)	0.55691 (15)	0.29311 (9)	0.0229 (4)
H9A	0.3458	0.5144	0.2843	0.027*
H9B	0.2868	0.6025	0.3348	0.027*
C10	0.25045 (16)	0.63354 (15)	0.23103 (9)	0.0258 (4)
H10A	0.3144	0.6878	0.2270	0.031*
H10B	0.1771	0.6745	0.2387	0.031*
C11	0.24125 (17)	0.56687 (17)	0.16328 (10)	0.0305 (4)
H11A	0.3155	0.5278	0.1547	0.037*
H11B	0.2273	0.6178	0.1242	0.037*
C12	0.10630 (14)	0.43289 (14)	0.25731 (9)	0.0194 (3)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Cl1	0.0465 (3)	0.0281 (3)	0.0379 (3)	0.0064 (2)	0.0101 (2)	-0.0076 (2)
S1	0.0275 (3)	0.0321 (3)	0.0220 (2)	-0.00856 (19)	-0.00050 (18)	-0.00137 (18)
S2	0.0221 (2)	0.0233 (2)	0.0301 (2)	-0.00669 (17)	-0.00089 (18)	-0.00064 (17)

supplementary materials

N1	0.0159 (7)	0.0190 (7)	0.0222 (7)	-0.0002 (5)	-0.0002 (6)	-0.0020 (6)
C1	0.0184 (8)	0.0250 (8)	0.0236 (9)	-0.0008 (7)	-0.0032 (7)	-0.0004 (7)
C2	0.0185 (8)	0.0219 (8)	0.0327 (10)	0.0015 (7)	0.0007 (7)	0.0034 (7)
C3	0.0264 (9)	0.0197 (8)	0.0282 (9)	-0.0021 (7)	0.0084 (7)	-0.0006 (7)
C4	0.0304 (10)	0.0293 (9)	0.0192 (8)	-0.0021 (8)	0.0022 (7)	0.0013 (7)
C5	0.0253 (9)	0.0246 (8)	0.0220 (8)	0.0020 (7)	-0.0003 (7)	0.0042 (7)
C6	0.0161 (8)	0.0198 (8)	0.0215 (8)	-0.0023 (6)	0.0012 (7)	0.0028 (7)
C7	0.0213 (9)	0.0204 (8)	0.0215 (8)	-0.0001 (7)	-0.0020 (7)	0.0007 (7)
C8	0.0342 (11)	0.0283 (9)	0.0325 (10)	0.0106 (8)	-0.0070 (8)	-0.0031 (8)
C9	0.0159 (8)	0.0247 (8)	0.0280 (9)	-0.0043 (7)	0.0015 (7)	-0.0042 (7)
C10	0.0214 (9)	0.0206 (8)	0.0353 (10)	-0.0045 (7)	0.0038 (8)	-0.0005 (8)
C11	0.0269 (10)	0.0359 (10)	0.0287 (9)	-0.0090 (8)	0.0034 (8)	0.0026 (8)
C12	0.0172 (8)	0.0158 (7)	0.0253 (8)	0.0024 (6)	-0.0003 (7)	-0.0020 (7)

Geometric parameters (Å, °)

C11—C3	1.7425 (18)	C5—H5	0.9300
S1—C12	1.7422 (18)	C6—C7	1.515 (2)
S1—C11	1.8092 (19)	C7—C8	1.525 (2)
S2—C12	1.6875 (17)	C7—H7	0.9800
N1—C12	1.330 (2)	C8—H8A	0.9600
N1—C9	1.481 (2)	C8—H8B	0.9600
N1—C7	1.485 (2)	C8—H8C	0.9600
C1—C2	1.383 (2)	C9—C10	1.512 (3)
C1—C6	1.394 (2)	C9—H9A	0.9700
C1—H1	0.9300	C9—H9B	0.9700
C2—C3	1.381 (3)	C10—C11	1.514 (3)
C2—H2	0.9300	C10—H10A	0.9700
C3—C4	1.379 (3)	C10—H10B	0.9700
C4—C5	1.389 (2)	C11—H11A	0.9700
C4—H4	0.9300	C11—H11B	0.9700
C5—C6	1.391 (2)		
C12—S1—C11	105.85 (8)	C7—C8—H8A	109.5
C12—N1—C9	124.50 (14)	C7—C8—H8B	109.5
C12—N1—C7	120.47 (14)	H8A—C8—H8B	109.5
C9—N1—C7	114.98 (13)	C7—C8—H8C	109.5
C2—C1—C6	121.49 (16)	H8A—C8—H8C	109.5
C2—C1—H1	119.3	H8B—C8—H8C	109.5
C6—C1—H1	119.3	N1—C9—C10	113.05 (14)
C3—C2—C1	119.00 (16)	N1—C9—H9A	109.0
C3—C2—H2	120.5	C10—C9—H9A	109.0
C1—C2—H2	120.5	N1—C9—H9B	109.0
C4—C3—C2	121.23 (16)	C10—C9—H9B	109.0
C4—C3—C11	119.37 (15)	H9A—C9—H9B	107.8
C2—C3—C11	119.36 (14)	C9—C10—C11	110.98 (15)
C3—C4—C5	119.06 (17)	C9—C10—H10A	109.4
C3—C4—H4	120.5	C11—C10—H10A	109.4
C5—C4—H4	120.5	C9—C10—H10B	109.4
C4—C5—C6	121.22 (16)	C11—C10—H10B	109.4

C4—C5—H5	119.4	H10A—C10—H10B	108.0
C6—C5—H5	119.4	C10—C11—S1	110.72 (12)
C5—C6—C1	117.99 (16)	C10—C11—H11A	109.5
C5—C6—C7	122.29 (15)	S1—C11—H11A	109.5
C1—C6—C7	119.64 (15)	C10—C11—H11B	109.5
N1—C7—C6	109.69 (13)	S1—C11—H11B	109.5
N1—C7—C8	110.24 (14)	H11A—C11—H11B	108.1
C6—C7—C8	115.75 (14)	N1—C12—S2	125.51 (13)
N1—C7—H7	106.9	N1—C12—S1	122.66 (13)
C6—C7—H7	106.9	S2—C12—S1	111.83 (9)
C8—C7—H7	106.9		

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C9—H9A \cdots S2 ⁱ	0.97	2.85	3.773 (2)	158
C10—H10B \cdots S2 ⁱⁱ	0.97	2.77	3.701 (2)	160

Symmetry codes: (i) $x+1/2, y, -z+1/2$; (ii) $-x, y+1/2, -z+1/2$.

Fig. 1

