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Key indicators

Single-crystal X-ray study T = 120 KMean $\sigma(C-C) = 0.003 \text{ Å}$ R factor = 0.044 wR factor = 0.147 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.

2-(*N*,*N*-Dimethylaminomethyl)-4-hydroxy-4-(4-chlorophenyl)thiazoline

The structure of the title compound, $C_{12}H_{15}ClN_2OS$, comprises a twisted molecule that forms dimers in the solid state *via* the O-H···N(thiazole) hydrogen-bonding interaction. The phenyl ring is twisted by 88.65 (9)° with respect to the mean plane of the thiazoline ring (largest ring deviation <0.05 Å). No close contacts are observed involving the amino N atom, but one of the *N*-methyl C atoms is 3.367 (3) Å from an adjacent Cl atom, although none of the *N*-methyl H atoms are appropriately positioned to form a C-H···Cl close contact. Received 22 May 2002 Accepted 28 May 2002 Online 8 June 2002



Experimental

The title compound was obtained from Key Organics Ltd and crystals were grown from an ethanol solution.

Crystal data	
$C_{12}H_{15}CIN_2OS$	Z = 2
$M_r = 270.77$	$D_x = 1.361 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 6.1186(8) Å	Cell parameters from 25743
b = 9.3865(9) Å	reflections
c = 11.977 (2) Å	$\theta = 1.0-27.5^{\circ}$
$\alpha = 103.473 \ (4)^{\circ}$	$\mu = 0.43 \text{ mm}^{-1}$
$\beta = 97.274 \ (5)^{\circ}$	T = 120 (2) K
$\gamma = 93.32 \ (1)^{\circ}$	Needle, colourless
$V = 660.8 (1) \text{ Å}^3$	$0.34 \times 0.07 \times 0.04 \text{ mm}$
Data collection	
Bruker-Nonius KappaCCD area-	2846 independent reflections

Bruker–Nonius KappaCCD areadetector diffractometer φ and ω scans Absorption correction: multi-scan (*SORTAV*; Blessing, 1995) $T_{\rm min} = 0.867, T_{\rm max} = 0.983$ 8739 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.147$ S = 1.012846 reflections 160 parameters 2206 reflections with $I > 2\sigma(I)$ $R_{int} = 0.042$ $\theta_{max} = 27.5^{\circ}$ $h = -7 \rightarrow 7$ $k = -12 \rightarrow 12$ $l = -15 \rightarrow 15$ H atoms treated by a mixture of independent and constrained

refinement $w = 1/[\sigma^2(F_o^2) + (0.1P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\text{max}} < 0.001$ $\Delta\rho_{\text{max}} = 0.34 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{\text{min}} = -0.39 \text{ e} \text{ Å}^{-3}$

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Table 1		
Hydrogen-bonding geometry	(Å,	°).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$O41-H41\cdots N3^i$	0.91 (3)	1.86 (3)	2.767 (2)	176 (3)
$C21 - H22 \cdot \cdot \cdot O41^{ii}$	0.99	2.38	3.294 (2)	153
$C42 - H42 \cdots O41$	0.95	2.43	2.793 (3)	102

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

All H atoms were included in the refinement, at calculated positions, as riding models, with C–H set to 0.93 (Ar–H) and 0.96 Å (CH₃), except for the O–H atom, which was located in a difference synthesis, and whose positional and displacement parameters were refined.

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

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Figure 1

The molecular configuration and atom-numbering scheme for the title compound, showing 50% probability ellipsoids.

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