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Key indicators

Single-crystal X-ray study T = 93 K Mean σ (C–C) = 0.003 Å R factor = 0.034 wR factor = 0.096 Data-to-parameter ratio = 12.5

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

2,9-Dichloroquino[2,3-b]acridine-7,14(5H,12H)dithione dimethylformamide disolvate

The title compound, $C_{20}H_{10}Cl_2N_2S_2 \cdot 2C_3H_7NO$, is a dimethylformamide (DMF) disolvate of DTQ-Cl, which is a thionated pigment derivative of 2,9-dichloroquinacridone. The DTQ-Cl molecule is centrosymmetric and entirely planar. Two DMF molecules are hydrogen-bonded to DTQ-Cl, through the NH group of DTQ-Cl and the O atom of DMF. The molecular plane of DMF is twisted with respect to the skeleton of DTQ-Cl by 21.4 (1)°.

Comment

The title compound, (I), is a DMF disolvate of DTQ-Cl, which is a thionated derivative of 2,9-dichloroquinacridone known as an industrially important red pigment (Herbst & Hunger, 1997). We have previously reported that thionation of unsubstituted quinacridone brings about an intense near-IR absorption that can be utilized for laser printers, as well as optical disks based on GaAsAl laser diodes (Mizuguchi et al., 1992, 1994). DTQ-Cl has, therefore, been synthesized in the expectation that it exhibits even better performance and stability for electronic applications. Since optical absorption in the solid state depends largely on intermolecular interactions, X-ray crystal structure analysis of DTQ-Cl is crucial. We obtained three types of solvated crystals: one was isolated from a DMF solution (present study) and the other two were from a single solution in dimethylacetamide (Senju et al., 2005; Hoki et al., 2005). This report deals with the title dimethylformamide solvate, DTQ-Cl·2DMF, (I).



Fig. 1 shows an *ORTEPIII* (Burnett & Johnson, 1996) plot of (I) including the two solvent molecules. The DTQ-Cl molecule is centrosymmetric and entirely planar, as characterized by the mean deviation of 0.006 Å from the leastsquares plane (C1–C10/N1).

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Figure 1

A view of the molecular structure of (I), showing 50% probability displacement ellipsoids. Unlabelled atoms are related to labelled atoms by the symmetry operator (-x, 2 - y, -z).

There are N-H···O intermolecular hydrogen bonds between the NH group of DTQ-Cl and the O atom of DMF. As shown in Table 2, the N, H and O atoms are perfectly aligned, with an N···O distance of 2.769 (2) Å, indicating that this hydrogen bond is quite strong. The molecular plane of DMF is twisted with respect to the skeleton of DTQ-Cl by 21.4 (1)°. DTQ-Cl molecules are stacked along the b axis in a herringbone fashion.

Experimental

DTQ-Cl was synthesized by thionation of commercially available 2,9dichloroquinacridone, using Lawesson's reagent (Rochat et al., 1988). Crystals of (I) were grown by gradual cooling of a dimethylformamide solution prepared at about 420 K.

Crystal data

$C_{20}H_{10}Cl_2N_2S_2 \cdot 2C_3H_7NO$	$D_x = 1.461 \text{ Mg m}^{-3}$		
$M_r = 559.53$	Cu $K\alpha$ radiation		
Monoclinic, $C2/c$	Cell parameters from 9508		
a = 33.257 (4) Å	reflections		
b = 4.2682 (6) Å	$\theta = 3.2-68.3^{\circ}$		
c = 21.589 (3) Å	$\mu = 4.10 \text{ mm}^{-1}$		
$\beta = 123.872 \ (7)^{\circ}$	T = 93.1 K		
V = 2544.4 (6) Å ³	Needle, dark green		
Z = 4	$0.50 \times 0.10 \times 0.05 \ \mathrm{mm}$		
Data collection			
Rigaku R-AXIS RAPID-F imaging	2123 independent reflections		
plate diffractometer	1947 reflections with $F^2 > 2\sigma(F^2)$		
ωscans	$R_{\rm int} = 0.028$		
Absorption correction: multi-scan	$\theta_{\rm max} = 68.2^{\circ}$		
(ABSCOR: Higashi, 1995)	$h = -40 \rightarrow 40$		

 $k = -4 \rightarrow 4$

 $l = -24 \rightarrow 25$

Refinement

refinement

Refinement on F^2
$R[F^2 > 2\sigma(F^2)] = 0.034$
$wR(F^2) = 0.096$
S = 1.10
2123 reflections
170 parameters
H atoms treated by a mixture of
independent and constrained

 $w = 1/[\sigma^2(F_0^2) + (0.0528P)^2]$ + 3.2176*P*] where $P = (F_0^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\rm max} < 0.001$ $\Delta\rho_{\rm max} = 0.40 \text{ e } \text{\AA}^{-3}_{\circ}$ $\Delta \rho_{\rm min} = -0.31 \text{ e } \text{\AA}^{-3}$

Table 1 Selected geometric parameters (Å, °).

Cl1-C2	1.746 (2)	C4-C5	1.418 (3)
S1-C7	1.663 (3)	C5-C6	1.411 (3)
N1-C5	1.357 (2)	C9-C10	1.393 (2)
N1-C9	1.372 (2)	C8-C9	1.418 (3)
C1-C2	1.371 (3)	C8-C7	1.462 (2)
C1-C6	1.413 (2)	C8-C10 ⁱ	1.394 (3)
C2-C3	1.404 (4)	C7-C8	1.462 (2)
C3-C4	1.364 (3)	C7-C6	1.458 (3)
Cl1-C2-C1	119.9 (2)	C1-C6-C7	121.4 (2)
Cl1-C2-C3	118.4 (1)	C1-C6-C5	118.2 (2)
S1-C7-C8	121.7 (2)	C2-C3-C4	119.4 (2)
S1-C7-C6	122.9 (1)	C3-C4-C5	120.2 (2)
N1-C5-C6	121.2 (2)	C4-C5-C6	120.3 (2)
N1-C5-C4	118.5 (2)	C5-C6-C7	120.4 (2)
C9-N1-C5	122.2 (2)	C9-C8-C7	121.0 (2)
N1-C9-C8	119.8 (2)	C10-C9-C8	120.8 (2)
N1-C9-C10	119.4 (2)	C9-C10-C8i	121.6 (2)
C1-C2-C3	121.7 (2)	C8-C7-C6	115.4 (2)
C6-C1-C2	120.1 (2)	$C10^{i} - C8 - C7$	121.4 (2)

Symmetry code: (i) -x, 2 - y, -z.

Fable 2	
Hydrogen-bond geometry (Å,	°).

$D - \mathbf{H} \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
N1−H1 <i>N</i> …O1	0.82 (2)	1.95 (3)	2.769 (2)	180 (3)

The position of the imide H atom was found in a difference Fourier map and refined freely along with an isotropic displacement parameter. Methyl H atoms were constrained to an ideal geometry, with C-H = 0.98 Å and $U_{iso}(H) = 1.2U_{eq}(C)$, but each group was allowed to rotate freely about its C-C bond. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C-H = 0.95 Å and $U_{iso}(H) = 1.2U_{eq}(C)$.

Data collection: PROCESS-AUTO (Rigaku, 1998); cell refinement: PROCESS-AUTO; data reduction: CrystalStructure (Rigaku/ MSC and Rigaku, 2004); program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: ORTEPIII (Burnett & Johnson, 1996); software used to prepare material for publication: CrystalStructure.

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 $T_{\min} = 0.280, \ T_{\max} = 0.815$

11 564 measured reflections

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