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## Structure Reports

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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 $=0.094$
$w R$ factor $=0.251$
Data-to-parameter ratio $=13.3$

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.
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## Ethyl 2-(2-chloro-1,4-dihydro-1,4-dioxo-naphthalen-3-ylamino)-4-phenylthiazole-5-carboxylate

The structure of the title compound, $\mathrm{C}_{22} \mathrm{H}_{15} \mathrm{ClN}_{2} \mathrm{O}_{4} \mathrm{~S}$, comprises non-planar molecules that form a one-dimensional hydrogen-bonded chain via a single $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ interaction, which runs parallel to the $b$ axis. The dihedral angle between the thiazole and quinone rings is $50.43(7)^{\circ}$ and the dihedral angle between the thiazole and the phenyl rings is $52.4(1)^{\circ}$.

## Comment

The title compound, (I), was prepared with the intention of merging two separate studies that we have recently undertaken. One study involved the synthesis and structural properties of 2-substituted 3-chloro-1,4-naphthoquinones (Lynch \& McClenaghan, 2002; 2003), while the other involved 2aminothiazoles. From the latter study came the structure of the thiazole derivative used to prepare (I), viz. ethyl 2-amino-4-phenylthiazole-5-carboxylate (Lynch \& McClenaghan, 2000). By bringing together the two series of molecules, we are interested in examining the combined structural aspects of the resultant covalently linked products, especially considering the forced proximity of one $\mathrm{N}-\mathrm{H}$ hydrogen-bond donor with five hydrogen-bond acceptors (viz. two O atoms, one N atom, one Cl atom and one S atom). The structure of (I) comprises non-planar molecules, the dihedral angle between the thiazole and quinone rings being 50.43 (7) ${ }^{\circ}$ and the dihedral angle between the thiazole and phenyl rings being 52.4 (1) ${ }^{\circ}$. The equivalent dihedral angle in the parent thiazole molecule is 42.41 (6) ${ }^{\circ}$.


Molecules of (I) form a one-dimensional hydrogen-bonded chain via a single $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ interaction [graph set $C(6)$; Etter, 1990], which runs parallel to the $b$ axis; hydrogen-bonding geometry is given in Table 1. A close contact $\mathrm{C} 25-$ $\mathrm{H} 25 \cdots \mathrm{O} 21^{\mathrm{i}}\left[\mathrm{C} \cdots \mathrm{O}^{\mathrm{i}}=3.165\right.$ (3) $\AA, \mathrm{H} \cdots \mathrm{O}^{\mathrm{i}}=2.22 \AA$ and $\mathrm{C}-$ $\mathrm{H} \cdots \mathrm{O}^{\mathrm{i}}=172^{\circ}$; symmetry code: (i) $\left.x, 1+y, z\right]$ exists adjacent to the $\mathrm{N}-\mathrm{H} \cdots \mathrm{O}$ interaction and thus completes an $R_{2}^{2}(10)$ graph-set motif.

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## Experimental

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

## Crystal data

$\mathrm{C}_{22} \mathrm{H}_{15} \mathrm{ClN}_{2} \mathrm{O}_{4} \mathrm{~S}$
$M_{r}=438.87$
Monoclinic, $P 2_{d} / c$
$a=19.191(5) \AA$
$b=7.719(2) \AA$
$c=12.640(3) \AA$
$\beta=94.845(18)^{\circ}$
$V=1865.6(8) \AA^{3}$
$Z=4$

$$
\begin{aligned}
& D_{x}=1.563 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation } \\
& \text { Cell parameters from } 4585 \\
& \quad \text { reflections } \\
& \theta=2.9-27.5^{\circ} \\
& \mu=0.35 \mathrm{~mm}^{-1} \\
& T=120(2) \mathrm{K} \\
& \text { Plate, orange } \\
& 0.18 \times 0.14 \times 0.02 \mathrm{~mm}
\end{aligned}
$$

## Data collection

Nonius KappaCCD diffractometer $\varphi$ and $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 2003) $T_{\text {min }}=0.939, T_{\text {max }}=0.993$
34099 measured reflections
3672 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.068$
$\theta_{\text {max }}=26.0^{\circ}$
$h=-23 \rightarrow 23$
$k=-9 \rightarrow 9$
$l=-15 \rightarrow 15$
3678 independent reflections

## Refinement

Refinement on $F^{2}$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.094$
$w R\left(F^{2}\right)=0.251$
$S=1.14$
3678 reflections
276 parameters
H atoms treated by a mixture of independent and constrained refinement

$$
\begin{gathered}
\begin{array}{c}
w=1 /\left[\sigma^{2}\left(F_{o}{ }^{2}\right)+(0.0933 P)^{2}\right. \\
\quad+12.2716 P] \\
\text { where } P=\left(F_{o}{ }^{2}+2 F_{c}^{2}\right) / 3 \\
(\Delta / \sigma)_{\max }<0.001 \\
\Delta \rho_{\max }=0.66 \mathrm{e}^{-3} \AA^{-3} \\
\Delta \rho_{\min }=
\end{array}-0.62 \mathrm{e}^{-3}
\end{gathered}
$$

Table 1
Hydrogen-bonding geometry $\left(\AA,^{\circ}\right)$.

| $D-\mathrm{H} \cdots A$ | $D-\mathrm{H}$ | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :--- | :--- | :--- | :--- | :--- |
| $\mathrm{~N} 21-\mathrm{H} 21 \cdots \mathrm{O} 24^{\mathrm{i}}$ | $0.86(2)$ | $2.27(2)$ | $3.066(3)$ | $154(2)$ |

Symmetry code: (i) $x, y-1, z$.

The amino H atom was located in a difference Fourier synthesis and its positional parameters were refined. Other H atoms were included in the refinement at calculated positions in the riding-model approximation, with $\mathrm{C}-\mathrm{H}$ distances of 0.95 (aromatic H atoms), 0.98 $\left(\mathrm{CH}_{3} \mathrm{H}\right.$ atoms $)$ and $0.99 \AA\left(\mathrm{CH}_{2} \mathrm{H}\right.$ atoms $)$. The isotropic displacement parameters for all H atoms were set equal to $1.25 U_{\text {eq }}$ of the


Figure 1
The molecular configuration and atom-numbering scheme for (I). Displacement ellipsoids are drawn at the $50 \%$ probability level and $H$ atoms are drawn as spheres of arbitrary radius.
carrier atom. The high $R$ value in this structure was a direct consequence of poor data from poor-quality twinned crystals; the nonmerohedral twinning was refined as two components with ratio 0.5207 (8):0.4793 (8).

Data collection: COLLECT (Hooft, 1998); cell refinement: DENZO (Otwinowski \& Minor, 1997) and COLLECT; data reduction: $D E N Z O$ and COLLECT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97.

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