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Daniel E. Lynch^a* and Ian McClenaghan^b†

^aSchool of Natural and Environmental Sciences, Coventry University, Coventry CV1 5FB, England, and ^bSpa Contract Synthesis, School of Natural and Environmental Sciences, Coventry University, Coventry CV1 5FB, England

+ E-mail: 106355.1670@CompuServe.com.

Correspondence e-mail: apx106@coventry.ac.uk

Key indicators

Single-crystal X-ray study T = 150 KMean $\sigma(C-C) = 0.004 \text{ Å}$ R factor = 0.061 wR factor = 0.170 Data-to-parameter ratio = 16.6

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

(E)-1-(2-Thienyl)ethanone 8-quinolylhydrozone

The structure of the title compound, $C_{15}H_{13}N_3S$, comprises planar molecules. The dihedral angle between the quinoline and thiophene rings is 2.6 (2)°.

Comment

Two distinct crystal colours (orange and yellow) were observed in the bulk following crystallization of the title compound, (I). The molecular structures from both crystal types were determined and proved to be identical, as were the melting points. Both structures refined to approximately the same *R* value and the structure obtained from the orange crystals is reported here. The cell for the yellow crystals was determined as *a* = 10.1194 (9), *b* = 14.251 (1), *c* = 18.136 (2) Å and *V* = 2615.4 (4) Å³. The origin of the difference in colour is currently unknown and cannot be simply explained by crystal morphology because both crystal types were indistinguishable in size and shape. However, we cannot discard the possibility of two different rates of crystallization, which we are currently investigating.



Experimental

Converted data

The title compound, (I), was prepared by Spa Contract Synthesis. Crystals were grown from an aqueous solution.

$C_{15}H_{13}N_{3}S$ $M_{r} = 267.34$ Orthorhombic, <i>Pbca</i> $a_{r} = 10.148(2) \text{ Å}$	Mo $K\alpha$ radiation Cell parameters from 46446 reflections $\theta = 1.0-27.5^\circ$
$ \begin{array}{l} a = 10.140 (2) \ A^{*} \\ b = 14.251 (3) \ A^{*} \\ c = 18.103 (4) \ A^{*} \\ V = 2618.1 (9) \ A^{3} \\ Z = 8 \\ D_{x} = 1.357 \ \text{Mg m}^{-3} \end{array} $	$\mu = 0.24 \text{ mm}^{-1}$ T = 150 (2) K Plate, orange $0.30 \times 0.24 \times 0.02 \text{ mm}$
Data collection Enraf-Nonius KappaCCD area- detector diffractometer φ and ω scans Absorption correction: multi-scan (SORTAV; Blessing, 1995) $T_{min} = 0.933, T_{max} = 0.995$ 20 334 measured reflections	2947 independent reflections 1498 reflections with $I > 2\sigma(I)$ $R_{int} = 0.121$ $\theta_{max} = 27.5^{\circ}$ $h = -11 \rightarrow 13$ $k = -18 \rightarrow 18$ $I = -23 \rightarrow 21$

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Refinement	
Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.061$ $wR(F^2) = 0.170$ S = 0.98 2947 reflections 177 parameters	H atoms treated by a mixture of independent and constrained refinement $w = 1/[\sigma^2(F_o^2) + (0.0867P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.32 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{min} = -0.33 \text{ e } \text{\AA}^{-3}$
Table 1 Hydrogen-bonding geometry (Å, °).	

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$		
N11-H11···N1	0.97 (4)	2.36 (3)	2.675 (4)	98 (2)		

All H atoms were included in the refinement at calculated positions as riding models with C–H set to 0.95 (Ar-H) and 0.98 Å (CH₃), except for the amine H atom, which was located on difference syntheses and for which both positional and displacement parameters were refined. The higher than expected R_{int} value (>0.10) may be due to a slight misalignment of the crystal plate on the diffractometer with respect to the beam.

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); software used to prepare material for publication: *SHELXL*97.

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Figure 1

The molecular configuration and atom numbering scheme for (I), showing 30% probability ellipsoids.

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