

(2E)-1-(2,5-Dimethylthiophen-3-yl)-3-(3-nitrophenyl)prop-2-en-1-one

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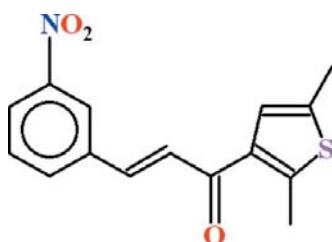
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Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(C-C) = 0.004$ Å; R factor = 0.047; wR factor = 0.117; data-to-parameter ratio = 13.5.

In the title compound, $C_{15}H_{13}NO_3S$, the benzene ring and the five-membered heterocyclic ring are oriented at a dihedral angle of $12.00(6)^\circ$. In the crystal, C–H···O interactions generate two types of cyclic motifs, $R_2^2(14)$ and $R_2^2(26)$, connecting the molecules into tapes extending along [101]. In addition, there are π – π stacking interactions between the benzene and thiophene rings with centroid-centroid distances of $3.7263(14)$ and $3.7487(14)$ Å.

Related literature

For the synthesis of similar compounds, see: Asiri & Khan (2010, 2011); Kalirajan *et al.* (2009); Patil *et al.* (2009); Sarojini *et al.* (2006). For related structures and background references, see: Asiri *et al.* (2010a,b). For graph-set notation, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$C_{15}H_{13}NO_3S$
 $M_r = 287.32$
Monoclinic, $P2_1/c$

$a = 7.3802(5)$ Å
 $b = 13.7973(9)$ Å
 $c = 13.4638(8)$ Å

$\beta = 96.997(3)^\circ$
 $V = 1360.77(15)$ Å³
 $Z = 4$
Mo $K\alpha$ radiation

$\mu = 0.24$ mm⁻¹
 $T = 296$ K
 $0.25 \times 0.22 \times 0.20$ mm

Data collection

Bruker KAPPA APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2005)
 $T_{\min} = 0.945$, $T_{\max} = 0.955$

10732 measured reflections
2466 independent reflections
1493 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.051$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.047$
 $wR(F^2) = 0.117$
 $S = 1.03$
2466 reflections

183 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.17$ e Å⁻³
 $\Delta\rho_{\min} = -0.25$ e Å⁻³

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

D–H···A	D–H	H···A	D···A	D–H···A
C6–H6···O3 ⁱ	0.93	2.46	3.373 (3)	168
C15–H15B···O2 ⁱⁱ	0.96	2.59	3.339 (4)	135

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

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997) and *PLATON* (Spek, 2009); software used to prepare material for publication: *WinGX* (Farrugia, 1999) and *PLATON*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2432).

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