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3,3'-Bis(4-nitrophenyl)-1,1'-(*p*-phenylene)dithiourea dimethyl sulfoxide disolvate

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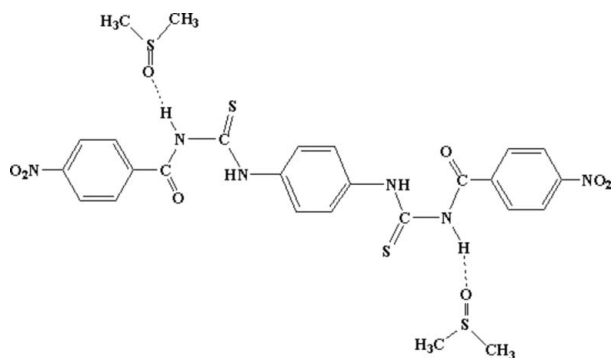
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 Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.006$ Å; R factor = 0.069; wR factor = 0.200; data-to-parameter ratio = 13.5.

The asymmetric unit of the title compound, $\text{C}_{22}\text{H}_{16}\text{N}_6\text{O}_6\text{S}_2 \cdot 2\text{C}_2\text{H}_6\text{OS}$, consists of one half-molecule of the centrosymmetric thiourea derivative and one molecule of dimethyl sulfoxide (DMSO). The carbonyl group forms an intramolecular hydrogen bond with the NH group, creating a six-membered (C—N—C—N—H \cdots O) ring. Two other N—H \cdots O hydrogen bonds link one molecule of the thiourea to two molecules of DMSO.

Related literature

For related literature, see: Burrows *et al.* (1997); Dong *et al.* (2006, 2007); Foss *et al.* (2004); Valdés-Martínez *et al.* (2000, 2004); Zhang *et al.* (2006); Huang *et al.* (2006).



Experimental

Crystal data

 $\text{C}_{22}\text{H}_{16}\text{N}_6\text{O}_6\text{S}_2 \cdot 2\text{C}_2\text{H}_6\text{OS}$
 $M_r = 680.78$

 Monoclinic, $P2_1/c$
 $a = 11.6949$ (18) Å
 $b = 6.6916$ (11) Å
 $c = 20.449$ (2) Å
 $\beta = 106.353$ (2)°
 $V = 1535.5$ (4) Å³
 $Z = 2$
 Mo $K\alpha$ radiation
 $\mu = 0.37$ mm⁻¹
 $T = 298$ (2) K
 $0.33 \times 0.17 \times 0.11$ mm

Data collection

 Bruker SMART CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.888$, $T_{\max} = 0.961$

 7318 measured reflections
 2684 independent reflections
 1547 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.097$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.069$
 $wR(F^2) = 0.199$
 $S = 0.96$
 2684 reflections

 199 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.44$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.43$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{O4}$	0.86	2.09	2.942 (5)	169
$\text{N2}-\text{H2}\cdots\text{O1}$	0.86	1.84	2.579 (5)	143

Data collection: SMART (Siemens, 1996); cell refinement: SAINT (Siemens, 1996); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 2008); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008); molecular graphics: SHELXTL (Sheldrick, 2008); software used to prepare material for publication: SHELXTL.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: FL2195).

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supporting information

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3,3'-Bis(4-nitrophenyl)-1,1'-(*p*-phenylene)dithiourea dimethyl sulfoxide disolvate

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S1. Comment

Thiourea and its derivatives are of interest to us because of their varied biological activity as well as their ability to form strong H bonds as both donors and acceptors (Valdés-Martínez, *et al.*, 2000; Jesus, *et al.*, 2004, Burrows *et al.*, 1997), and their tendency to coordinate with metal ions (Huang, *et al.*, 2006; Foss, *et al.*, 2004). In recent years, thioureas have been recognized as important neutral receptors because of their anion recognition properties (Zhang, *et al.*, 2006) and for their ability to easily form intramolecular hydrogen bonds such as between the benzoyl (CO) and the N—H group of acylthioureas (Dong *et al.*, 2006). In continuation of our previous studies on the synthesis and structural characterization of *N*-benzoyl-*N'*-(3-pyridyl)thiourea (II) (Dong, *et al.*, 2006) and *N,N'*-(1,6-hexamethylene)-bis(benzoylthiourea) (Dong, *et al.*, 2007), a novel bisbenzoylthiourea which crystallized as a dimethyl sulfoxide disolvate (I) has now been synthesized and structurally characterized.

The crystal structure of (I) is built up by one *N,N'*-(*p*-phenyl)-bis(*p*-nitro)benzoylthiourea molecule and two dimethyl sulfoxide solvent molecules. The carbonyl group of the thiourea forms an intramolecular hydrogen bond with the N—H group to form a six-membered (C/N/C/N/H/O) ring. The C=O bond length at 1.225 (5) Å is longer than the average C=O bond length (1.200 Å). This is most likely due to the intramolecular hydrogen bonding which is similar to the situation found in the structure of (II) (Dong, *et al.*, 2006).

There are also N—H···O hydrogen bonds between N1 of the thiourea and the O atoms of the DMSO S=O groups which, together with other intermolecular C—H···O and C—H···S interactions, stabilize the three-dimensional structure of (I).

S2. Experimental

(*p*-Nitro)benzoyl chloride (1.86 g, 10 mmol) was reacted with ammonium thiocyanate (1.14 g, 15 mmol) in CH₂Cl₂ (25 ml) solution under solid-liquid phase transfer catalysis, using polyethylene glycol-400 (0.18 g) as the catalyst, to give the corresponding (*p*-nitro)benzoyl isothiocyanate under stirring at the room temperature. This was followed by slow addition of 15 ml CH₂Cl₂ solution dissolved *p*-phenylenediamine (1.60 g, 0.01 mmol). The corresponding yellow compound precipitated immediately. The product was filtered, washed with water and CH₂Cl₂, dried, and recrystallized from THF to give the titled thiourea. Yield, 72.6%. m. p. 243 - 244 °C. Anal. Calc. for C₂₂H₁₆N₆O₆S₂ (%): C, 50.38; H, 3.07; N, 16.02. Found: C, 50.27; H, 3.15; N, 15.99. Selected IR data (cm⁻¹, KBr pellet): 3341, 3191 (ν NH), 1675 (ν C=O), 1152 (ν C=S). ¹H NMR (400 MHz, DMSO-d₆, δ, p.p.m.): 8.05 (d, J = 17.2 Hz, 4H, ArH), 8.19 (d, J = 8.2 Hz, 4H, ArH), 8.34 (dd, J = 17.2, 7.2 Hz, 4H, ArH), 11.98 (s, 2H, NH), 12.42 (s, 2H, NH).

A DMSO solution of the thiourea was placed in a hexane atmosphere, after about one week, along with diffusion of hexane into the DMSO solution, yellow needle-shaped single crystals suitable for X-ray crystallographic analysis were obtained.

S3. Refinement

Non-H atoms were refined anisotropically. H atoms were treated as riding atoms with distances C—H = 0.96(CH₂), or 0.93 Å (CH), O—H = 0.86 Å, and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.5U_{\text{eq}}(\text{O})$.

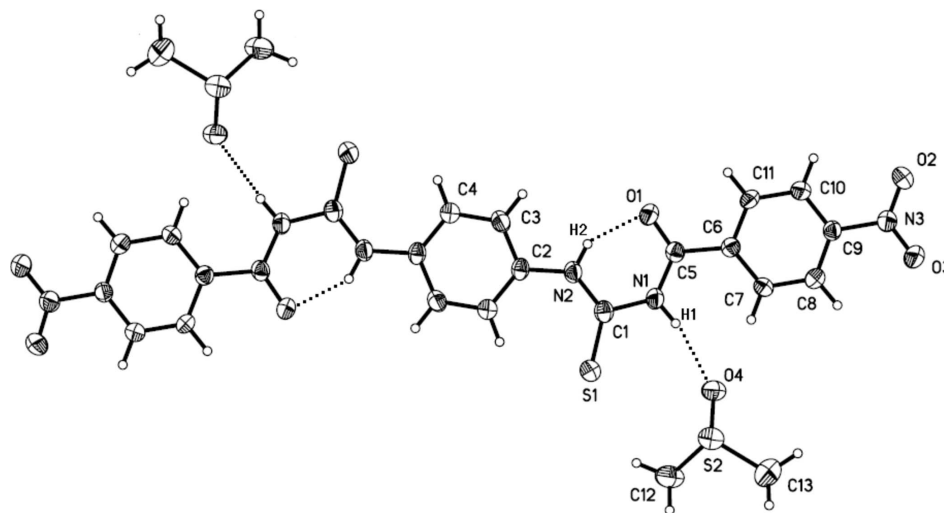


Figure 1

Molecule structure of (I) with the atom numbering. Displacement ellipsoids for non-H atoms are drawn at the 30% probability level.

3,3'-Bis(4-nitrophenyl)-1,1'-(p-phenylene)dithiourea dimethyl sulfoxide disolvate

Crystal data

$\text{C}_{22}\text{H}_{16}\text{N}_6\text{O}_6\text{S}_2 \cdot 2\text{C}_2\text{H}_6\text{OS}$

$M_r = 680.78$

Monoclinic, $P2_1/c$

$a = 11.6949$ (18) Å

$b = 6.6916$ (11) Å

$c = 20.449$ (2) Å

$\beta = 106.353$ (2)°

$V = 1535.5$ (4) Å³

$Z = 2$

$F(000) = 708$

$D_x = 1.472$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 1517 reflections

$\theta = 3.3\text{--}25.3^\circ$

$\mu = 0.37$ mm⁻¹

$T = 298$ K

Needle-shaped, yellow

$0.33 \times 0.17 \times 0.11$ mm

Data collection

Bruker SMART CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Sheldrick, 1996)

$T_{\text{min}} = 0.888$, $T_{\text{max}} = 0.961$

7318 measured reflections

2684 independent reflections

1547 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.097$

$\theta_{\text{max}} = 25.0^\circ$, $\theta_{\text{min}} = 1.8^\circ$

$h = -11 \rightarrow 13$

$k = -7 \rightarrow 7$

$l = -24 \rightarrow 24$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.069$

$wR(F^2) = 0.199$

$S = 0.97$

2684 reflections

199 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods
Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.1071P)^2]$$

$$\text{where } P = (F_o^2 + 2F_c^2)/3$$

$$(\Delta/\sigma)_{\max} < 0.001$$

$$\Delta\rho_{\max} = 0.44 \text{ e } \text{\AA}^{-3}$$

$$\Delta\rho_{\min} = -0.43 \text{ e } \text{\AA}^{-3}$$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
N1	0.7828 (3)	0.3522 (5)	0.51603 (16)	0.0505 (8)
H1	0.7673	0.2724	0.5454	0.061*
N2	0.8719 (3)	0.6436 (5)	0.49516 (16)	0.0507 (9)
H2	0.8343	0.6109	0.4541	0.061*
N3	0.4791 (3)	-0.4240 (5)	0.36972 (18)	0.0565 (9)
O1	0.7411 (3)	0.4141 (5)	0.40259 (14)	0.0614 (8)
O2	0.4552 (3)	-0.4678 (5)	0.31005 (16)	0.0733 (10)
O3	0.4566 (3)	-0.5286 (5)	0.41241 (17)	0.0776 (10)
O4	0.7382 (3)	0.1230 (6)	0.62917 (14)	0.0782 (10)
S1	0.91836 (14)	0.5258 (2)	0.62543 (6)	0.0858 (6)
S2	0.82807 (12)	0.0691 (2)	0.69426 (6)	0.0711 (5)
C1	0.8572 (4)	0.5173 (6)	0.5429 (2)	0.0502 (10)
C2	0.9391 (3)	0.8218 (6)	0.5010 (2)	0.0454 (10)
C3	0.9492 (4)	0.9013 (7)	0.4410 (2)	0.0557 (11)
H3	0.9141	0.8344	0.4004	0.067*
C4	1.0090 (4)	1.0755 (7)	0.4389 (2)	0.0562 (11)
H4	1.0150	1.1251	0.3975	0.067*
C5	0.7326 (3)	0.3047 (6)	0.44901 (19)	0.0455 (10)
C6	0.6642 (3)	0.1137 (6)	0.43250 (18)	0.0443 (9)
C7	0.6443 (4)	-0.0164 (6)	0.4804 (2)	0.0501 (10)
H7	0.6723	0.0141	0.5265	0.060*
C8	0.5824 (4)	-0.1933 (7)	0.4596 (2)	0.0548 (11)
H8	0.5674	-0.2810	0.4915	0.066*
C9	0.5441 (3)	-0.2359 (6)	0.39199 (19)	0.0461 (10)
C10	0.5631 (4)	-0.1108 (7)	0.3439 (2)	0.0600 (12)
H10	0.5363	-0.1438	0.2979	0.072*
C11	0.6225 (4)	0.0652 (7)	0.3643 (2)	0.0592 (12)
H11	0.6349	0.1531	0.3318	0.071*
C12	0.8181 (5)	0.2507 (9)	0.7540 (2)	0.0870 (17)

H12A	0.8448	0.3769	0.7414	0.131*
H12B	0.8671	0.2124	0.7983	0.131*
H12C	0.7368	0.2624	0.7550	0.131*
C13	0.7650 (6)	-0.1309 (11)	0.7298 (3)	0.120 (2)
H13A	0.6928	-0.0866	0.7388	0.180*
H13B	0.8206	-0.1736	0.7716	0.180*
H13C	0.7477	-0.2405	0.6983	0.180*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
N1	0.050 (2)	0.046 (2)	0.0538 (19)	-0.0108 (17)	0.0109 (15)	0.0002 (17)
N2	0.0460 (19)	0.047 (2)	0.0551 (19)	-0.0107 (16)	0.0077 (15)	-0.0008 (17)
N3	0.064 (2)	0.048 (2)	0.055 (2)	-0.0041 (18)	0.0123 (18)	0.0011 (18)
O1	0.0683 (19)	0.057 (2)	0.0569 (17)	-0.0188 (15)	0.0140 (14)	0.0064 (15)
O2	0.094 (2)	0.065 (2)	0.0564 (19)	-0.0207 (18)	0.0137 (17)	-0.0127 (16)
O3	0.100 (3)	0.064 (2)	0.064 (2)	-0.0294 (19)	0.0153 (18)	0.0067 (17)
O4	0.090 (2)	0.090 (3)	0.0495 (18)	-0.021 (2)	0.0117 (16)	0.0026 (17)
S1	0.1133 (12)	0.0803 (10)	0.0544 (8)	-0.0440 (9)	0.0080 (7)	-0.0012 (6)
S2	0.0732 (9)	0.0804 (10)	0.0590 (7)	0.0026 (7)	0.0174 (6)	-0.0010 (6)
C1	0.043 (2)	0.048 (3)	0.059 (3)	-0.0024 (19)	0.0118 (19)	-0.002 (2)
C2	0.035 (2)	0.040 (2)	0.060 (2)	-0.0015 (17)	0.0101 (17)	0.0030 (19)
C3	0.059 (3)	0.049 (3)	0.054 (2)	-0.009 (2)	0.0082 (19)	-0.007 (2)
C4	0.064 (3)	0.055 (3)	0.050 (2)	-0.011 (2)	0.016 (2)	-0.002 (2)
C5	0.035 (2)	0.048 (3)	0.050 (2)	-0.0030 (18)	0.0056 (17)	0.001 (2)
C6	0.040 (2)	0.047 (2)	0.046 (2)	-0.0016 (18)	0.0121 (17)	0.0012 (18)
C7	0.051 (2)	0.053 (3)	0.042 (2)	-0.006 (2)	0.0060 (18)	-0.0026 (19)
C8	0.062 (3)	0.052 (3)	0.053 (2)	-0.008 (2)	0.021 (2)	0.004 (2)
C9	0.046 (2)	0.042 (2)	0.047 (2)	-0.0039 (18)	0.0067 (17)	0.0017 (19)
C10	0.073 (3)	0.061 (3)	0.042 (2)	-0.018 (2)	0.011 (2)	-0.001 (2)
C11	0.073 (3)	0.060 (3)	0.046 (2)	-0.023 (2)	0.017 (2)	0.008 (2)
C12	0.101 (4)	0.097 (4)	0.059 (3)	0.013 (3)	0.016 (3)	-0.001 (3)
C13	0.158 (6)	0.091 (5)	0.100 (4)	-0.022 (5)	0.020 (4)	0.024 (4)

Geometric parameters (Å, °)

N1—C5	1.368 (4)	C4—H4	0.9300
N1—C1	1.418 (5)	C5—C6	1.495 (6)
N1—H1	0.8600	C6—C7	1.379 (6)
N2—C1	1.338 (5)	C6—C11	1.380 (5)
N2—C2	1.414 (5)	C7—C8	1.390 (6)
N2—H2	0.8600	C7—H7	0.9300
N3—O3	1.204 (4)	C8—C9	1.358 (5)
N3—O2	1.209 (4)	C8—H8	0.9300
N3—C9	1.475 (5)	C9—C10	1.356 (6)
O1—C5	1.225 (5)	C10—C11	1.371 (6)
O4—S2	1.490 (3)	C10—H10	0.9300
S1—C1	1.638 (4)	C11—H11	0.9300

S2—C12	1.750 (5)	C12—H12A	0.9600
S2—C13	1.779 (6)	C12—H12B	0.9600
C2—C3	1.372 (5)	C12—H12C	0.9600
C2—C4 ⁱ	1.389 (5)	C13—H13A	0.9600
C3—C4	1.367 (6)	C13—H13B	0.9600
C3—H3	0.9300	C13—H13C	0.9600
C4—C2 ⁱ	1.389 (5)		
C5—N1—C1	127.8 (3)	C11—C6—C5	116.4 (3)
C5—N1—H1	116.1	C6—C7—C8	119.9 (4)
C1—N1—H1	116.1	C6—C7—H7	120.1
C1—N2—C2	130.8 (3)	C8—C7—H7	120.1
C1—N2—H2	114.6	C9—C8—C7	119.0 (4)
C2—N2—H2	114.6	C9—C8—H8	120.5
O3—N3—O2	123.7 (4)	C7—C8—H8	120.5
O3—N3—C9	118.1 (3)	C10—C9—C8	122.3 (4)
O2—N3—C9	118.1 (4)	C10—C9—N3	118.6 (3)
O4—S2—C12	106.6 (2)	C8—C9—N3	119.2 (4)
O4—S2—C13	106.1 (2)	C9—C10—C11	118.8 (4)
C12—S2—C13	96.9 (3)	C9—C10—H10	120.6
N2—C1—N1	113.6 (3)	C11—C10—H10	120.6
N2—C1—S1	128.4 (3)	C10—C11—C6	121.0 (4)
N1—C1—S1	117.9 (3)	C10—C11—H11	119.5
C3—C2—C4 ⁱ	118.2 (4)	C6—C11—H11	119.5
C3—C2—N2	115.9 (4)	S2—C12—H12A	109.5
C4 ⁱ —C2—N2	125.8 (4)	S2—C12—H12B	109.5
C4—C3—C2	122.1 (4)	H12A—C12—H12B	109.5
C4—C3—H3	118.9	S2—C12—H12C	109.5
C2—C3—H3	118.9	H12A—C12—H12C	109.5
C3—C4—C2 ⁱ	119.6 (4)	H12B—C12—H12C	109.5
C3—C4—H4	120.2	S2—C13—H13A	109.5
C2 ⁱ —C4—H4	120.2	S2—C13—H13B	109.5
O1—C5—N1	122.2 (4)	H13A—C13—H13B	109.5
O1—C5—C6	119.4 (3)	S2—C13—H13C	109.5
N1—C5—C6	118.4 (4)	H13A—C13—H13C	109.5
C7—C6—C11	119.1 (4)	H13B—C13—H13C	109.5
C7—C6—C5	124.5 (3)		
C2—N2—C1—N1	-179.4 (4)	C11—C6—C7—C8	-0.3 (6)
C2—N2—C1—S1	-2.6 (7)	C5—C6—C7—C8	-178.1 (4)
C5—N1—C1—N2	4.2 (6)	C6—C7—C8—C9	1.0 (6)
C5—N1—C1—S1	-173.0 (3)	C7—C8—C9—C10	-0.7 (7)
C1—N2—C2—C3	170.9 (4)	C7—C8—C9—N3	179.6 (4)
C1—N2—C2—C4 ⁱ	-11.5 (7)	O3—N3—C9—C10	-175.8 (4)
C4 ⁱ —C2—C3—C4	0.6 (7)	O2—N3—C9—C10	6.8 (6)
N2—C2—C3—C4	178.4 (4)	O3—N3—C9—C8	4.0 (6)
C2—C3—C4—C2 ⁱ	-0.6 (7)	O2—N3—C9—C8	-173.4 (4)
C1—N1—C5—O1	-5.3 (6)	C8—C9—C10—C11	-0.3 (7)

C1—N1—C5—C6	175.5 (4)	N3—C9—C10—C11	179.4 (4)
O1—C5—C6—C7	-177.0 (4)	C9—C10—C11—C6	1.1 (7)
N1—C5—C6—C7	2.3 (6)	C7—C6—C11—C10	-0.7 (7)
O1—C5—C6—C11	5.2 (6)	C5—C6—C11—C10	177.2 (4)
N1—C5—C6—C11	-175.6 (4)		

Symmetry code: (i) $-x+2, -y+2, -z+1$.

Hydrogen-bond geometry (Å, °)

<i>D—H...A</i>	<i>D—H</i>	<i>H...A</i>	<i>D...A</i>	<i>D—H...A</i>
N1—H1...O4	0.86	2.09	2.942 (5)	169
N2—H2...O1	0.86	1.84	2.579 (5)	143