



Crystal structures of three *N,N,N'*-trisubstituted thioureas for reactivity-controlled nanocrystal synthesis

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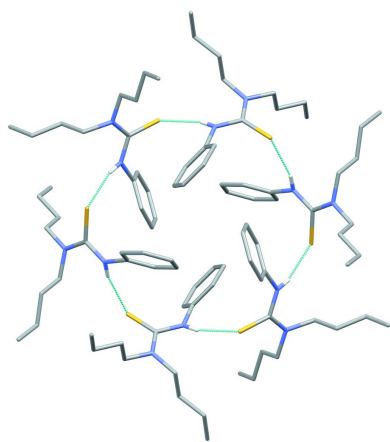
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The synthesis and single-crystal X-ray structures of three *N,N,N'*-trisubstituted thioureas are reported, namely *N,N,N'*-tribenzylthiourea, C₂₂H₂₂N₂S (**1**), *N*-methyl-*N,N'*-diphenylthiourea, C₁₄H₁₄N₂S (**2**), and *N,N*-di-*n*-butyl-*N'*-phenylthiourea, C₁₅H₂₄N₂S (**3**). The influence of the different substituents on the thioureas is clear from the delocalization of the thiourea C–N and C=S bonds, while the crystal structures show infinite chains of *N,N,N'*-tribenzylthiourea (**1**), hydrogen-bonded pairs of *N*-methyl-*N,N'*-diphenylthiourea (**2**) and hexamer ring assemblies of *N,N*-di-*n*-butyl-*N'*-phenylthiourea (**3**) molecules. The above-mentioned compounds were synthesized *via* a mild, general procedure, readily accessible precursors and with a high yield, providing straightforward access to a whole library of thioureas.

1. Chemical context

To control the size of colloidal nanocrystals, many traditional methods terminate the synthesis during the nanocrystal growth at the desired size. However, this leads to a lower yield, higher size dispersity, and it is difficult to get a good reproducibility (Owen *et al.*, 2010; Abe *et al.*, 2012, 2013). Therefore, Owen *et al.* suggest a new method that uses a library of substituted thioureas, whose substitution pattern tunes their conversion reactivity (Hendricks *et al.*, 2015; Hamachi *et al.*, 2017). By this, the nanocrystal concentration can be adjusted and the desired nanocrystal size can be obtained at full conversion, with a high degree of consistency. This control is obtained by varying the substitution pattern of the thiourea, and thus the conversion reactivity (Hens, 2015). This can be understood from the fact that the conversion reactivity is influenced by the number of substituents, and their electronic and steric properties. The conversion rate, *i.e.* reactivity, decreases as the number of substituents increases, or by replacing electron-withdrawing with electron-donating groups (*e.g.* substituting aryl for alkyl substituents). These thioureas are synthesized *via* a one-step click reaction between isothiocyanates and primary or secondary amines (Hendricks *et al.*, 2015). In addition, they have a long shelf-life and are air-stable after synthesis (Hendricks *et al.*, 2015). An additional advantage of these precursors is that the starting reagents are relatively cheap and widely commercially available, in large quantities. When added to a hot solution of metal oleate, such as lead, cadmium, zinc, *etc.*, this results in the formation of highly reproducible, monodisperse, homogeneously capped



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metal sulfide nanocrystals at a full yield (Hendricks *et al.*, 2015; Hamachi *et al.*, 2017; Dhaene *et al.*, 2019).



- 1:** $R_1 = -\text{Bn}$; $R_2 = -\text{Bn}$; $R_3 = -\text{Bn}$
2: $R_1 = -\text{Ph}$; $R_2 = -\text{Ph}$; $R_3 = -\text{Me}$
3: $R_1 = -\text{Ph}$; $R_2 = -\text{Bu}$; $R_3 = -\text{Bu}$

Herein, we report the single-crystal X-ray structural analysis of the following trisubstituted thioureas: *N,N,N'*-tribenzylthiourea (**1**), *N*-methyl-*N,N'*-diphenylthiourea (**2**), and *N*-phenyl-*N',N'*-di-*n*-butylthiourea (**3**), prepared *via* a simple, straightforward synthesis method making use of readily commercially available compounds, to a high purity (> 99%) and with a high yield (> 75%).

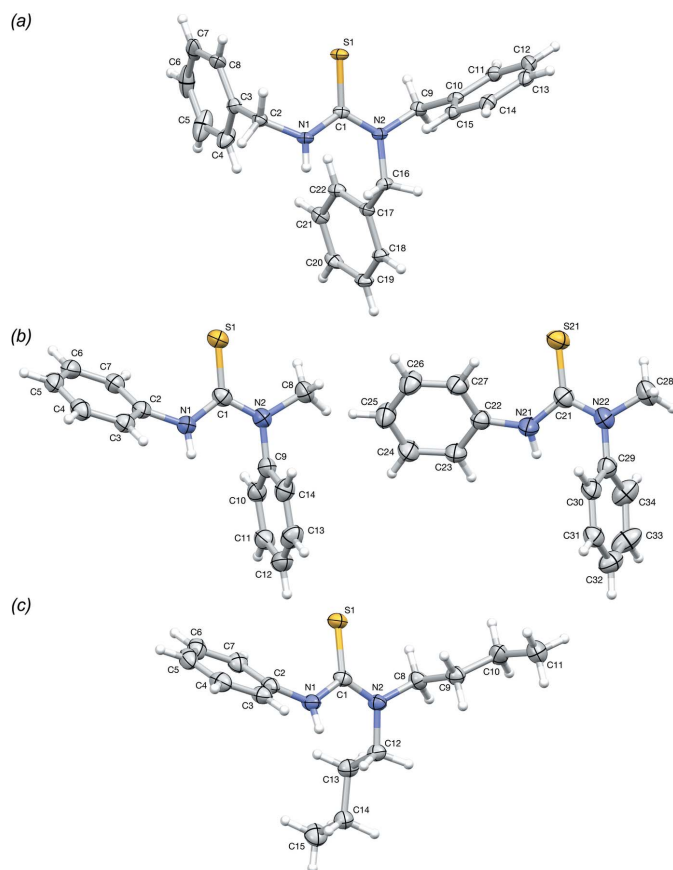


Figure 1
Molecular structures of (a) **1**, (b) **2** and (c) **3**, showing thermal displacement ellipsoids drawn at the 50% probability level and the atom-labelling scheme for the non-hydrogen atoms. For **2**, both molecules of the asymmetric unit are shown.

2. Structural commentary

Compound **1** crystallizes in the centrosymmetric monoclinic space group $P2_1/c$, with the asymmetric unit consisting of one *N,N,N'*-tribenzylthiourea molecule. On the one hand, the secondary amine benzyl ring (C3–C8) is found to be almost completely parallel to one of the tertiary amine benzyl rings (C17–C22), subtending a dihedral angle of $8.92(8)^\circ$ between the best planes through the two benzene rings. On the other hand, the two tertiary amine benzyl rings (C10–C15 and C17–C22) are highly twisted to each other, with a dihedral angle of $76.96(7)^\circ$ between the best planes through the two benzene rings (Fig. 1*a*). The N1–C1 and C1–N2 bond distances are 1.3419 (18) and 1.3569 (18) Å, respectively, while the C1=S1 (double) bond distance is 1.6905 (14) Å.

Compound **2** crystallizes in the centrosymmetric triclinic space group $P\bar{1}$, with two *N*-methyl-*N,N'*-diphenylthiourea molecules in the asymmetric unit. The secondary and tertiary amine phenyl rings (C2–C7, C9–C14 and C22–C27, C29–C34, for the first and second molecules, respectively) subtend a dihedral angle of $69.39(9)$ and $75.70(10)^\circ$, respectively, between the best planes through the two phenyl rings (Fig. 1*b*). The N1–C1 and C1–N2 bond distances are 1.359 (2) and 1.352 (3) Å, for molecule 1, while the respective N21–C21 and C21–N22 bond distances are 1.367 (2) and 1.345 (3) Å, for molecule 2. The C1=S1 and C21=S22 (double) bond distances are 1.6835 (17) and 1.6798 (19) Å, for molecule 1 and 2, respectively.

The influence of the two phenyl substituents on the delocalization of the N1–C1, C1–N2 and C1=S1 bonds is clear, in comparison with the structure of **1**, *i.e.* the lone electron pair on N1/N21 is more delocalized towards the secondary amine phenyl ring substituent in **2**, leading to an increased N1–C1/N21–C22 distance of 1.359 (2)/1.367 (2) Å, which is even more pronounced for the second molecule in the asymmetric unit, because of higher planarity of the phenyl ring with the N–C(=S)–N plane (Fig. 2). However, the delocalization of

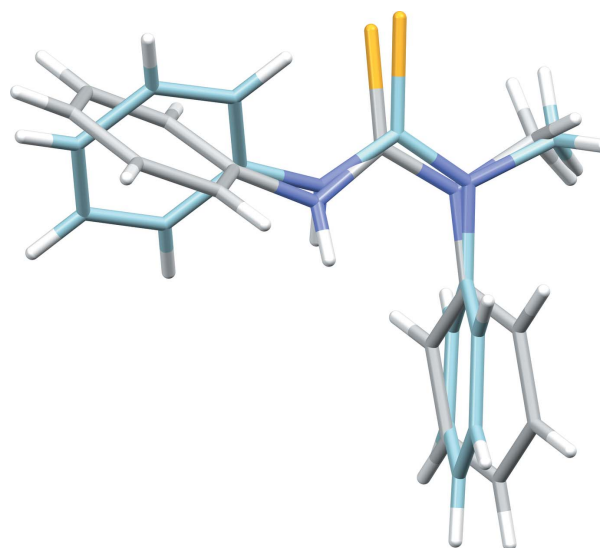


Figure 2
Fit of the first (grey) and second (light blue) molecule in the asymmetric unit of **2**, showing an r.m.s.d. of 1.174 Å.

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots S1^i$	0.86 (3)	2.47 (3)	3.2044 (13)	145 (3)

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

$N2/N22$ is less pronounced towards the tertiary amine phenyl ring, with a $C1-N2/C21-N22$ distance of 1.352 (3)/1.345 (3) Å, because of the latter phenyl ring being almost perpendicular to the central $N-C(=S)-N$ plane. As a consequence of the improved delocalization of $N1/N21$ in **2**, the $C1=S1/C21=S21$ bond length decreases slightly – although less significantly in the case of $C1=S1$ – to 1.6835 (17)/1.6798 (19) Å in comparison with **1**.

The structure of **3** has very recently been deposited with the Cambridge Structural Database (CSD) (refcode OYOSIH; Rahman *et al.*, 2021); however, the mentioned structure was determined at room temperature and showed disorder of both butyl substituents, as well as the presence of unknown solvent, which was treated by the SQUEEZE procedure in PLATON (Spek, 2015). Here, our reported structure was determined at 100 K and shows no signs of any kind of (solvent) disorder. The unknown solvate structure of Rahman *et al.* (2021) might be caused by the use of acetone as solvent and recrystallization by slow evaporation from EtOH, whereas we used toluene as solvent and recrystallized from a hot hexane:EtOH (10:1) mixture by slowly cooling down. Compound **3** crystallizes in the trigonal space group $R\bar{3}$, with one *N*-phenyl-*N,N'*-di-*n*-butylthiourea molecule in the asymmetric unit. The phenyl substituent on the secondary amine is twisted with respect to the central $N-C-S-N$ plane, with a $C1-N1-C2-C7$ torsion angle of 55.54 (16)°, while the two butyl substituents are found completely staggered (Fig. 1c). The $N1-C1$ and $C1-N2$ bond distances are 1.3594 (15) and 1.3432 (15) Å, respectively, while the $C1=S2$ (double) bond distance is 1.7004 (11) Å. The delocalization of $N1$ towards the secondary amine phenyl substituent is also noticed here, comparable to **2**, while there is minimal delocalization of $N2$ towards the butyl substituents, consequently showing the shortest $C1-N2$ and the longest $C1=S1$ distances.

3. Supramolecular features

Despite the presence of three benzyl moieties in the molecular structure of **1**, only weak $\pi-\pi$ interactions are present in the crystal packing, with rather large centroid–centroid distances ranging from 4.4279 (11) to 5.9248 (9) Å. However, clear intermolecular hydrogen bonds are found between the secondary amine $N1-H1$ hydrogen atoms and the thiourea $S1$ atoms [$N1-H1\cdots S1 = 2.47$ (3) Å; Table 1], linking the *N,N,N'*-tribenzylthiourea molecules into infinite chains along the [001] direction, and forming columnar arrangements through alternating orientations of the molecules (Fig. 3). Non-classical intramolecular hydrogen bonds can be noticed between methyl $C-H$ atoms of two benzyl groups and $S1$ atoms [$C2-H2A\cdots S1 = 2.70$ Å; $C9-H9A\cdots S1 = 2.60$ Å], as

Table 2
 Hydrogen-bond geometry (Å, °) for **2**.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots S21$	0.86 (3)	2.58 (3)	3.3360 (16)	148 (3)

well as between benzene ring $C-H$ atoms and tertiary amine $N2$ atoms [$C15-H15\cdots N2 = 2.51$ Å; $C22-H22\cdots N2 = 2.58$ Å]. Furthermore, several $C-H\cdots\pi$ contacts are observed in the range of 3.5419 (17)–3.8507 (19) Å, complementing the crystal packing.

Analogous to **1**, the presence of two phenyl substituents in the molecular structure of **2**, only leads to weak $\pi-\pi$ interactions present in the crystal packing, with rather large centroid–centroid distances ranging from 4.8431 (13) to 5.9503 (12) Å. However, in this case, intermolecular hydrogen bonds are formed between the two distinct molecules in the asymmetric unit, *i.e.* between the secondary amine $N1-H1$ hydrogen atom of the first molecule and the thiourea $S21$ atom

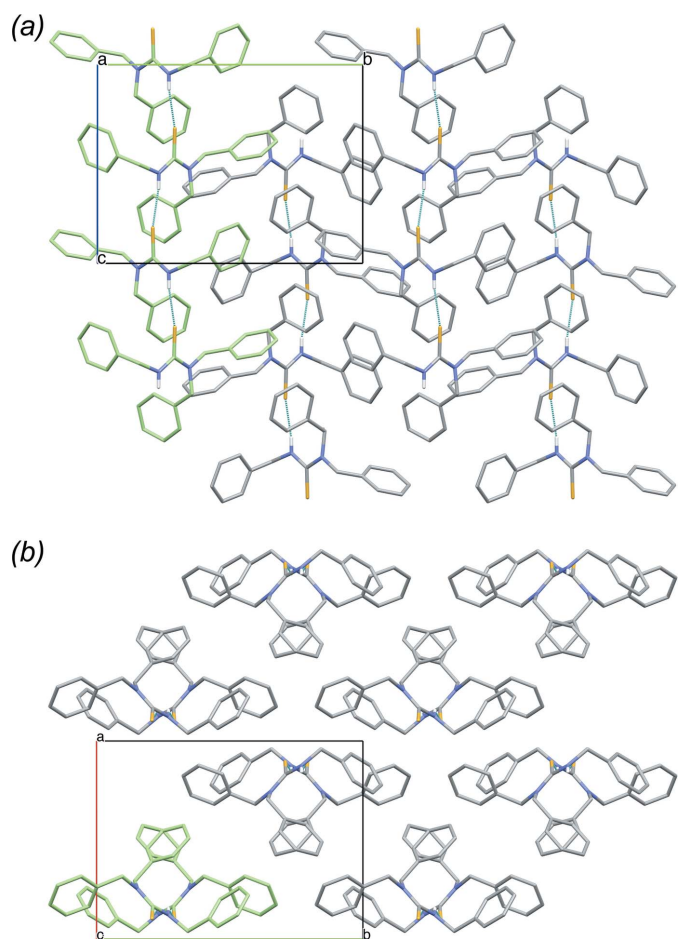


Figure 3
 Packing in the structure of **1**, (a) viewed down the a axis, showing the $N1-H1\cdots S1$ hydrogen bonds, linking the *N,N,N'*-tribenzylthiourea molecules into infinite chains along the [001] direction, and (b) viewed down the c axis, showing the columnar arrangement through alternating orientations of the molecules. A chain of four hydrogen-bonded molecules is highlighted (green). Hydrogen atoms (except involved in hydrogen bonds) are omitted for clarity.

Table 3
Hydrogen-bond geometry (Å, °) for **3**.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$N1-H1\cdots S1^i$	0.86 (2)	2.62 (2)	3.4656 (11)	167 (2)
$C12-H12A\cdots S1^i$	0.99	2.67	3.6588 (13)	174

Symmetry code: (i) $y - \frac{1}{3}, -x + y + \frac{1}{3}, -z + \frac{4}{3}$.

of the second molecule [$N1-H1\cdots S21 = 2.58$ (3) Å; Table 2], assembling the *N*-methyl-*N,N'*-diphenylthiourea molecules into hydrogen-bonded pairs (Fig. 4). Non-classical intramolecular hydrogen bonds can be noticed between the two methyl group C–H atoms, as well as phenyl ring C–H atoms, and S1/S21 atoms [$C8-H8B\cdots S1 = 2.65$ Å; $C28-H28B\cdots S21 = 2.58$ Å; $C27-H27\cdots S21 = 2.67$ Å]. Additionally, an intramolecular $C=S\cdots\pi$ contact is observed [$C21=S21\cdots Cg2 = 3.7115$ (11) Å; $Cg2$ is the centroid of the C9–C14 ring]. Furthermore, several C–H $\cdots\pi$ contacts are observed in the range of 3.518 (2)–3.800 (2) Å, complementing the crystal packing.

Analogous to **1** and **2**, for **3**, only one type of weak π – π interaction is present in the crystal packing, *i.e.* between symmetry-equivalent phenyl substituents, with a centroid–centroid distance of 4.9098 (10) Å. Intermolecular hydrogen bonds are formed between the secondary amine $N1-H1$ hydrogen atoms and the thiourea S1 atoms [$N1-H1\cdots S1 = 3.4656$ (11) Å; Table 3], leading to a hexamer ring assembly of molecules, around the threefold rotoinversion axes (Fig. 5). Non-classical intra- and intermolecular hydrogen bonds can be noticed between two butyl CH_2 groups and S1 [$C8-H8B\cdots S1 = 2.58$ Å; $C12-H12A\cdots S1^i$; symmetry code: (i) $-\frac{1}{3} + y, \frac{1}{3} - x + y, \frac{4}{3} - z$]. Only one C–H $\cdots\pi$ contact is observed [$C4-H4\cdots Cg1 = 3.6996$ (17) Å; $Cg1$ is the centroid of the C2–C7 ring].

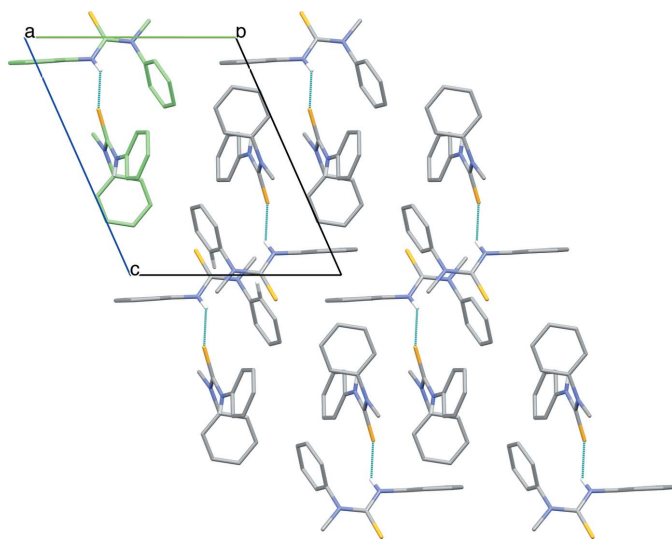


Figure 4
Packing in the structure of **2**, viewed down the *a* axis, showing the assembly of hydrogen-bonded pairs of molecules, with one pair highlighted (green). Hydrogen atoms (except involved in hydrogen bonds) are omitted for clarity.

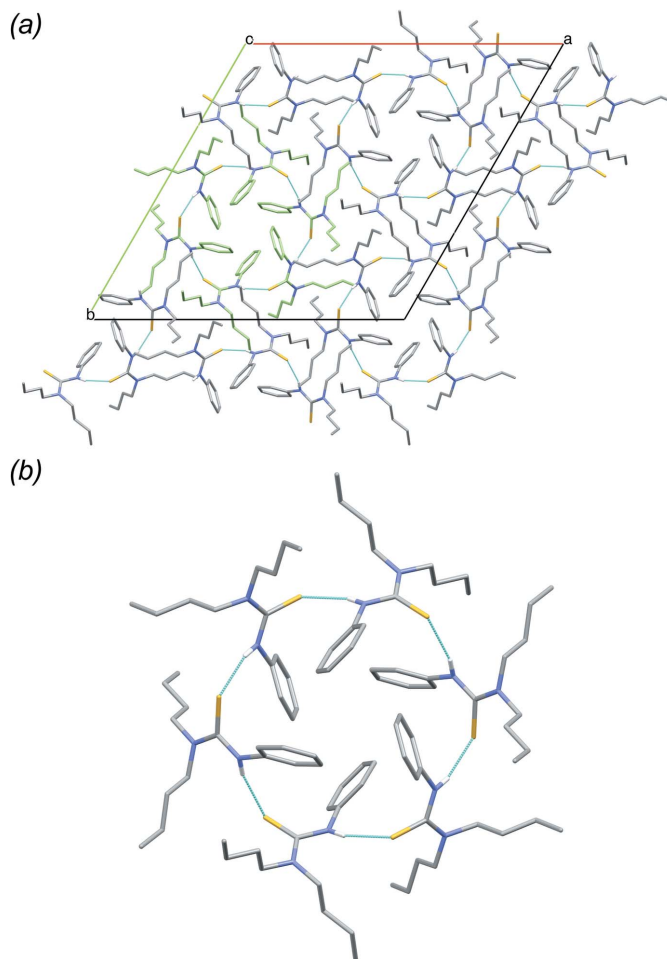


Figure 5
Packing in the structure of **3**, (a) viewed down the *c* axis, showing the hexamer ring assembly of molecules, around the threefold rotoinversion axes, with one hexamer highlighted (green). (b) Detail of one hydrogen-bonded hexamer ring assembly. Hydrogen atoms (except involved in hydrogen bonds) are omitted for clarity.

4. Database survey

A survey of compounds, closely related to **1**, **2** and **3**, deposited with the Cambridge Structural Database (CSD 2021.1, version 5.42, updates of September 2021; Groom *et al.*, 2016) resulted in ten other thiourea compounds, containing (substituted) benzyl/phenyl rings on the secondary amine and (substituted) benzyl/phenyl rings or alkyl groups on the tertiary amine, with refcodes HIFTIZ, HIFTOF, KUFQOS, KUFQOS01, KUFQOS02, POFJUR, QEMZOA, RAPNAA, RAQRAF and OYOSIH.

The structures with refcodes HIFTIX and HIFTOF are two unsymmetrical thiourea derivatives, 1,1-dimethyl-3-*o*-tolylthiourea and 1,1-diethyl-3-*o*-tolylthiourea (Ramnathan *et al.*, 1996), containing *o*-tolyl groups as secondary amine substituents, while KUFQOS (Zhao *et al.*, 2008), KUFQOS01 (Panda *et al.*, 2017) and KUFQOS02 (Bhide *et al.*, 2021) represent the same structure of 1,1-dimethyl-3-phenylthiourea. Halogen-substituted phenyl rings as secondary amine substituents are found for refcodes POFJUR and QEMZOA,

which represent isomorphic structures of 3-(2-bromo-4-chlorophenyl)-1,1-dimethylthiourea (El-Hiti *et al.*, 2014) and *N'*-(2-bromo-4-methylphenyl)-*N,N*-dimethylthiourea (El-Hiti *et al.*, 2018), respectively, while RAPNAA and RAQRAF represent structures of 3-(2-bromophenyl)-1,1-dimethylthiourea (El-Hiti *et al.*, 2017a) and 3-(4-chlorophenyl)-1,1-dimethylthiourea (El-Hiti *et al.*, 2017b), respectively.

In all the above-mentioned structures, N–H...S hydrogen bonds link the molecules into infinite chains, similar to **1**. This makes the reported structures of **2** and **3** unique in the sense that they show assemblies of hydrogen-bonded pairs and hexamer rings of molecules, respectively.

As previously mentioned, OYOSIH (Rahman *et al.*, 2021) represents the same structure as **3**, although determined at room temperature and showed disorder of both butyl substituents, as well as the presence of unknown solvent, which was treated by the SQUEEZE procedure in PLATON (Spek, 2015)

5. Synthesis and crystallization

General considerations All manipulations were performed in air. All chemicals were used as received. Phenyl isothiocyanate (97.0%) was purchased from Alfa Aesar. Chloroform-*d*₁ (stabilized with Ag, 99.8%D) was purchased from Carl Roth. Toluene (99.0%), acetonitrile (99.9%), *n*-hexane (99.0%), and abs. ethanol (99.8%) were purchased from Chem-Lab. Benzyl isothiocyanate (98.0%), dibenzylamine (97.0%), *N*-methylaniline (98.0%), and di-*n*-butylamine (99.5%) were purchased from Sigma-Aldrich. Dichloromethane-*d*₂ (99.8%D) was purchased from VWR. The thioureas were synthesized according to the procedure by Hendricks and Co-workers on a 30 mmol scale with the addition of a recrystallization step to purify the thiourea (Hendricks *et al.*, 2015; Hamachi *et al.*, 2017).

Synthesis of *N,N,N*-tribenzylthiourea (1**):** A 40 mL vial was loaded with benzyl isothiocyanate (4476.6 mg, 3.800 mL, 30 mmol, 1.0 eq.) in toluene (5 mL). To this, a solution of dibenzylamine (5918.4 mg, 5.800 mL, 30 mmol, 1.0 eq.) in toluene (5 mL) was added dropwise. The mixture was left to stir for 1 h at room temperature. Afterwards, the solvent was removed under reduced pressure, and the residual solid was recrystallized from hot acetonitrile which was cooled slowly (> 2 h) to room temperature and then to refrigerator temperature (275–281 K; > 2 h). The formed crystals were filtered off and extensively dried under dynamic vacuum to obtain white needle-like crystals (7.8 g, 75%), suitable for single-crystal X-ray diffraction analysis. ¹H NMR (400 MHz, CD₂Cl₂): δ 7.45–7.15 (*m*, 13H), δ 7.10–6.95 (*m*, 2H), δ 5.80 (*t*, *J* = 4.5 Hz, 1H), δ 5.00 (*s*, 4H), 4.80 (*d*, *J* = 2.6 Hz, 2H). ¹³C NMR (100 MHz, CDCl₃): δ 183.16, 137.78, 136.06, 129.17, 128.75, 128.03, 127.62, 127.19, 54.37, 50.79. **LC–MS** (API–ES) calculated for C₂₂H₂₃N₂S [*M*+H]⁺ 347.16, found 347.1.

Synthesis of *N*-methyl-*N,N*-diphenylthiourea (2**):** A 40 mL vial was loaded with phenyl isothiocyanate (4055.7 mg, 3.585 mL, 30 mmol, 1.0 eq.) in toluene (5 mL). To this, a solution of *N*-methylaniline (3214.5 mg, 3.250 mL, 30 mmol,

1.0 eq.) in toluene (5 mL) was added dropwise. The mixture was left to stir for 6 h at 323 K, since the reaction with aniline derivatives elapses more sluggishly. Afterwards, the solvent was removed under reduced pressure, and the residual solid was recrystallized from a hot hexane:ethanol (10:1) mixture which was cooled slowly (> 2 h) to room temperature and then to refrigerator temperature (275–281 K; > 2 h). The formed crystals were filtered off and extensively dried under dynamic vacuum to obtain white needle-like crystals (5.5 g, 76%), suitable for single-crystal X-ray diffraction analysis. ¹H NMR (400 MHz, CD₂Cl₂): δ 7.55–7.50 (*m*, 2H), δ 7.45–7.35 (*m*, 3H), δ 7.32–7.27 (*m*, 4H), δ 7.20–7.12 (*m*, 1H), δ 7.00 (*s*, 1H), δ 3.70 (*s*, 3H). ¹³C NMR (100 MHz, CDCl₃): δ 181.92, 143.52, 140.04, 131.05, 129.03, 128.76, 127.44, 126.21, 126.12, 43.73. **LC–MS** (API–ES) calculated for C₁₄H₁₅N₂S [*M*+H]⁺ 243.10, found 243.1.

Synthesis of *N,N*-di-*n*-butyl-*N'*-phenylthiourea (3**):** A 40 mL vial was loaded with phenyl isothiocyanate (4055.7 mg, 3.585 mL, 30 mmol, 1.0 eq.) in toluene (5 mL). To this, a solution of di-*n*-butylamine (3877.2 mg, 5.055 mL, 30 mmol, 1.0 eq.) in toluene (5 mL) was added dropwise. The mixture was left to stir for 1 h at room temperature. Afterwards, the solvent was removed under reduced pressure, and the residual solid was recrystallized from a hot hexane:ethanol (10:1) mixture which was cooled slowly (> 2 h) to room temperature and then to refrigerator temperature (275–281 K; > 2 h). The formed crystals were filtered off and extensively dried under dynamic vacuum to obtain white needle-like crystals (6.9 g, 87%), suitable for single-crystal X-ray diffraction analysis. ¹H NMR (400 MHz, CD₂Cl₂): δ 7.40–7.28 (*m*, 4H), δ 7.23–7.15 (*m*, 1H), δ 7.00 (*s*, 1H), δ 3.67 (*t*, *J* = 7.9 Hz, 4H), δ 1.71 (*quin*, *J* = 7.7 Hz, 4H), δ 1.38 (*six*, *J* = 7.9 Hz, 4H), δ 0.97 (*t*, *J* = 7.5 Hz, 6H). ¹³C NMR (100 MHz, CDCl₃): δ 181.56, 140.65, 128.85, 126.21, 125.84, 51.85, 29.95, 20.69, 14.06. **LC–MS** (API–ES) calc for C₁₅H₂₅N₂S [*M*+H]⁺ 265.17, found 265.2.

NMR spectroscopy. Nuclear Magnetic Resonance (NMR) spectra of the synthesized organics were recorded on a Bruker 400 MHz. Chemical shifts (δ) are given in ppm and the residual solvent peak was used as an internal standard (CDCl₃: δH = 7.24 ppm, δC = 77.06 ppm, CD₂Cl₂: δH = 5.32 ppm, δC = 53.84 ppm). The signal multiplicity is denoted as follows: *s* (singlet), *d* (doublet), *t* (triplet), *quad* (quadruplet), *quin* (quintet), *six* (sextet), *m* (multiplet). Coupling constants are reported in Hertz (Hz). All resonances were corrected prior to integration by subtracting a background from the measured intensity. ¹H, and ¹³C spectra were acquired using the standard pulse sequences from the Bruker library; zg30, and jmod (Attached Proton Test = APT), respectively.

Mass spectroscopy. Mass spectra (MS) were measured with an Agilent ESI single quadrupole detector type VL and an Agilent APCI single quadrupole detector type VL.

6. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 4. For all structures, the amine N–H hydrogen atoms could be located from a difference-Fourier

Table 4
Experimental details.

	1	2	3
Crystal data			
Chemical formula	C ₂₂ H ₂₂ N ₂ S	C ₁₄ H ₁₄ N ₂ S	C ₁₅ H ₂₄ N ₂ S
<i>M_r</i>	346.48	242.33	264.42
Crystal system, space group	Monoclinic, <i>P</i> 2 ₁ / <i>c</i>	Triclinic, <i>P</i> $\bar{1}$	Trigonal, <i>R</i> $\bar{3}$
Temperature (K)	100	100	100
<i>a</i> , <i>b</i> , <i>c</i> (Å)	11.2378 (4), 14.7792 (5), 11.3165 (5)	9.8379 (6), 10.8014 (6), 13.2328 (6)	25.5231 (3), 25.5231 (3), 12.6225 (2)
α , β , γ (°)	90, 102.042 (3), 90	65.913 (5), 87.752 (4), 84.059 (5)	90, 90, 120
<i>V</i> (Å ³)	1838.15 (12)	1276.82 (13)	7121.0 (2)
<i>Z</i>	4	4	18
Radiation type	Cu <i>K</i> α	Cu <i>K</i> α	Cu <i>K</i> α
μ (mm ⁻¹)	1.59	2.06	1.69
Crystal size (mm)	0.24 × 0.19 × 0.06	0.26 × 0.17 × 0.13	0.42 × 0.26 × 0.18
Data collection			
Diffractometer	SuperNova, Dual, Cu at home/ near, Atlas	SuperNova, Dual, Cu at home/ near, Atlas	SuperNova, Dual, Cu at home/ near, Atlas
Absorption correction	Gaussian (<i>CrysAlis PRO</i> ; Rigaku OD, 2019)	Gaussian (<i>CrysAlis PRO</i> ; Rigaku OD, 2019)	Gaussian (<i>CrysAlis PRO</i> ; Rigaku OD, 2019)
<i>T</i> _{min} , <i>T</i> _{max}	0.750, 1.000	0.687, 1.000	0.479, 1.000
No. of measured, independent and observed [<i>I</i> > 2 σ (<i>I</i>)] reflections	16515, 3589, 3256	12208, 4818, 4298	14921, 3151, 3004
<i>R</i> _{int}	0.060	0.028	0.021
(<i>sin</i> θ / λ) _{max} (Å ⁻¹)	0.624	0.623	0.622
Refinement			
<i>R</i> [<i>F</i> ² > 2 σ (<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.043, 0.123, 1.06	0.049, 0.142, 1.06	0.032, 0.087, 1.06
No. of reflections	3589	4818	3151
No. of parameters	229	315	168
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement	H atoms treated by a mixture of independent and constrained refinement	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\max}$, $\Delta\rho_{\min}$ (e Å ⁻³)	0.30, -0.31	0.58, -0.25	0.25, -0.18

Computer programs: *CrysAlis PRO* (Rigaku OD, 2019), *SHELXT2018/2* (Sheldrick, 2015a), *SHELXL2018/3* (Sheldrick, 2015b), *OLEX2* (Dolomanov *et al.*, 2009) and *Mercury* (Macrae *et al.*, 2020).

electron-density map, and were further refined with isotropic temperature factors fixed at 1.2 times *U*_{eq} of the parent atoms. All other hydrogen atoms were refined in the riding mode with isotropic temperature factors fixed at 1.2 times *U*_{eq} of the parent atoms (1.5 times for methyl groups).

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Crystal structures of three *N,N,N'*-trisubstituted thioureas for reactivity-controlled nanocrystal synthesis

Evert Dhaene, Isabel Van Driessche, Klaartje De Buysser and Kristof Van Hecke

Computing details

For all structures, data collection: *CrysAlis PRO* (Rigaku OD, 2019); cell refinement: *CrysAlis PRO* (Rigaku OD, 2019); data reduction: *CrysAlis PRO* (Rigaku OD, 2019); program(s) used to solve structure: *SHELXT2018/2* (Sheldrick, 2015a); program(s) used to refine structure: *SHELXL2018/3* (Sheldrick, 2015b); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *Mercury* (Macrae *et al.*, 2020).

N,N,N'-Tribenzylthiourea (1)

Crystal data

$C_{22}H_{22}N_2S$

$M_r = 346.48$

Monoclinic, $P2_1/c$

$a = 11.2378$ (4) Å

$b = 14.7792$ (5) Å

$c = 11.3165$ (5) Å

$\beta = 102.042$ (3)°

$V = 1838.15$ (12) Å³

$Z = 4$

$F(000) = 736$

$D_x = 1.252$ Mg m⁻³

Cu $K\alpha$ radiation, $\lambda = 1.54184$ Å

Cell parameters from 8187 reflections

$\theta = 4.0$ – 74.0 °

$\mu = 1.59$ mm⁻¹

$T = 100$ K

Plate, clear colourless

$0.24 \times 0.19 \times 0.06$ mm

Data collection

SuperNova, Dual, Cu at home/near, Atlas diffractometer

Radiation source: micro-focus sealed X-ray tube, SuperNova (Cu) X-ray Source

Mirror monochromator

Detector resolution: 10.4839 pixels mm⁻¹

ω scans

Absorption correction: gaussian (CrysAlisPro; Rigaku OD, 2019)

$T_{\min} = 0.750$, $T_{\max} = 1.000$

16515 measured reflections

3589 independent reflections

3256 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.060$

$\theta_{\max} = 74.2$ °, $\theta_{\min} = 4.0$ °

$h = -13 \rightarrow 13$

$k = -18 \rightarrow 18$

$l = -11 \rightarrow 13$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.043$

$wR(F^2) = 0.123$

$S = 1.06$

3589 reflections

229 parameters

0 restraints

Primary atom site location: dual

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0753P)^2 + 0.4416P]$

where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.30$ e Å⁻³

$\Delta\rho_{\min} = -0.31$ e Å⁻³

Absolute structure: -

Special details

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

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.10482 (3)	0.29165 (3)	0.32245 (3)	0.02217 (14)
N1	0.12474 (11)	0.23113 (8)	0.54565 (11)	0.0198 (3)
N2	0.25536 (10)	0.34685 (8)	0.52442 (11)	0.0191 (3)
C1	0.16633 (12)	0.28879 (9)	0.47205 (13)	0.0177 (3)
C2	0.04085 (13)	0.15697 (10)	0.50233 (13)	0.0209 (3)
H2A	-0.024231	0.179223	0.435803	0.025*
H2B	0.002213	0.136546	0.568717	0.025*
C3	0.10428 (12)	0.07750 (10)	0.45767 (14)	0.0215 (3)
C4	0.20059 (15)	0.03591 (12)	0.53478 (18)	0.0358 (4)
H4	0.226181	0.057328	0.615227	0.043*
C5	0.25996 (16)	-0.03682 (13)	0.4954 (2)	0.0519 (6)
H5	0.326738	-0.064616	0.548228	0.062*
H1	0.154 (3)	0.2304 (18)	0.622 (3)	0.062*
C6	0.22149 (18)	-0.06872 (13)	0.3786 (2)	0.0494 (6)
H6	0.262381	-0.118215	0.351143	0.059*
C7	0.1240 (2)	-0.02889 (12)	0.30214 (19)	0.0433 (5)
H7	0.096928	-0.051638	0.222546	0.052*
C8	0.06532 (16)	0.04457 (11)	0.34141 (15)	0.0299 (4)
H8	-0.001553	0.072197	0.288476	0.036*
C9	0.31744 (12)	0.40391 (10)	0.45137 (13)	0.0202 (3)
H9A	0.313944	0.374034	0.372425	0.024*
H9B	0.404170	0.408858	0.492068	0.024*
C10	0.26494 (12)	0.49838 (10)	0.42917 (13)	0.0190 (3)
C11	0.31752 (13)	0.55802 (10)	0.35960 (14)	0.0242 (3)
H11	0.384844	0.538845	0.327235	0.029*
C12	0.27248 (15)	0.64526 (11)	0.33705 (15)	0.0281 (4)
H12	0.308310	0.685134	0.288571	0.034*
C13	0.17537 (14)	0.67414 (11)	0.38522 (15)	0.0276 (4)
H13	0.144508	0.733810	0.370034	0.033*
C14	0.12353 (13)	0.61561 (10)	0.45559 (15)	0.0249 (3)
H14	0.057692	0.635536	0.489717	0.030*
C15	0.16720 (12)	0.52792 (10)	0.47662 (13)	0.0211 (3)
H15	0.130022	0.487852	0.523739	0.025*
C16	0.29405 (13)	0.35662 (10)	0.65564 (13)	0.0209 (3)
H16A	0.222424	0.347522	0.692503	0.025*
H16B	0.323109	0.419334	0.673830	0.025*
C17	0.39376 (12)	0.29173 (10)	0.71478 (14)	0.0195 (3)
C18	0.45690 (13)	0.31034 (11)	0.83213 (14)	0.0230 (3)
H18	0.438430	0.363493	0.871962	0.028*

C19	0.54635 (13)	0.25185 (11)	0.89097 (14)	0.0264 (3)
H19	0.589405	0.265389	0.970497	0.032*
C20	0.57320 (13)	0.17348 (11)	0.83395 (15)	0.0263 (3)
H20	0.634493	0.133350	0.874170	0.032*
C21	0.50997 (13)	0.15434 (11)	0.71817 (15)	0.0244 (3)
H21	0.527282	0.100429	0.679235	0.029*
C22	0.42111 (13)	0.21354 (10)	0.65824 (14)	0.0221 (3)
H22	0.379057	0.200265	0.578260	0.027*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0224 (2)	0.0317 (2)	0.0112 (2)	0.00067 (12)	0.00060 (14)	0.00133 (12)
N1	0.0220 (6)	0.0250 (6)	0.0118 (6)	0.0008 (5)	0.0020 (5)	0.0003 (5)
N2	0.0202 (6)	0.0220 (6)	0.0136 (6)	0.0015 (4)	0.0006 (4)	0.0009 (4)
C1	0.0162 (6)	0.0215 (7)	0.0153 (7)	0.0052 (5)	0.0028 (5)	0.0009 (5)
C2	0.0195 (7)	0.0251 (7)	0.0181 (7)	-0.0015 (5)	0.0041 (5)	0.0017 (6)
C3	0.0189 (7)	0.0240 (7)	0.0222 (8)	-0.0037 (5)	0.0057 (5)	0.0002 (6)
C4	0.0274 (8)	0.0291 (8)	0.0444 (11)	0.0016 (6)	-0.0073 (7)	-0.0044 (7)
C5	0.0243 (8)	0.0301 (10)	0.0956 (18)	0.0028 (7)	-0.0003 (9)	-0.0106 (10)
C6	0.0427 (11)	0.0253 (9)	0.0929 (18)	-0.0065 (7)	0.0434 (11)	-0.0113 (10)
C7	0.0719 (14)	0.0292 (9)	0.0394 (11)	-0.0185 (9)	0.0362 (10)	-0.0078 (8)
C8	0.0434 (9)	0.0277 (8)	0.0196 (8)	-0.0080 (7)	0.0090 (6)	0.0008 (6)
C9	0.0175 (6)	0.0251 (7)	0.0184 (7)	0.0009 (5)	0.0044 (5)	0.0013 (6)
C10	0.0178 (6)	0.0236 (7)	0.0141 (7)	-0.0007 (5)	-0.0002 (5)	-0.0016 (5)
C11	0.0242 (7)	0.0289 (8)	0.0194 (8)	-0.0031 (6)	0.0048 (5)	-0.0010 (6)
C12	0.0333 (8)	0.0272 (8)	0.0226 (9)	-0.0079 (6)	0.0029 (6)	0.0032 (6)
C13	0.0277 (8)	0.0216 (7)	0.0286 (9)	-0.0007 (6)	-0.0054 (6)	0.0010 (6)
C14	0.0190 (7)	0.0253 (8)	0.0283 (8)	0.0023 (5)	0.0001 (6)	-0.0020 (6)
C15	0.0180 (6)	0.0245 (7)	0.0196 (7)	-0.0009 (5)	0.0012 (5)	-0.0006 (6)
C16	0.0215 (7)	0.0251 (7)	0.0142 (8)	0.0027 (5)	-0.0003 (5)	-0.0012 (5)
C17	0.0166 (7)	0.0261 (7)	0.0156 (7)	-0.0013 (5)	0.0027 (5)	0.0031 (5)
C18	0.0226 (7)	0.0285 (7)	0.0167 (8)	-0.0028 (6)	0.0010 (5)	-0.0004 (6)
C19	0.0218 (7)	0.0373 (9)	0.0173 (8)	-0.0031 (6)	-0.0020 (5)	0.0038 (6)
C20	0.0169 (6)	0.0340 (8)	0.0264 (8)	0.0020 (6)	0.0012 (6)	0.0118 (7)
C21	0.0200 (7)	0.0279 (8)	0.0261 (9)	0.0030 (5)	0.0064 (6)	0.0027 (6)
C22	0.0192 (7)	0.0296 (8)	0.0167 (7)	0.0010 (5)	0.0019 (5)	0.0007 (6)

Geometric parameters (Å, °)

S1—C1	1.6905 (14)	C10—C15	1.390 (2)
N1—C1	1.3419 (19)	C11—H11	0.9500
N1—C2	1.4621 (18)	C11—C12	1.389 (2)
N1—H1	0.85 (3)	C12—H12	0.9500
N2—C1	1.3569 (18)	C12—C13	1.385 (3)
N2—C9	1.4566 (19)	C13—H13	0.9500
N2—C16	1.4648 (19)	C13—C14	1.384 (2)
C2—H2A	0.9900	C14—H14	0.9500

C2—H2B	0.9900	C14—C15	1.389 (2)
C2—C3	1.514 (2)	C15—H15	0.9500
C3—C4	1.384 (2)	C16—H16A	0.9900
C3—C8	1.385 (2)	C16—H16B	0.9900
C4—H4	0.9500	C16—C17	1.5199 (19)
C4—C5	1.387 (3)	C17—C18	1.396 (2)
C5—H5	0.9500	C17—C22	1.386 (2)
C5—C6	1.384 (3)	C18—H18	0.9500
C6—H6	0.9500	C18—C19	1.386 (2)
C6—C7	1.378 (3)	C19—H19	0.9500
C7—H7	0.9500	C19—C20	1.389 (2)
C7—C8	1.391 (3)	C20—H20	0.9500
C8—H8	0.9500	C20—C21	1.383 (2)
C9—H9A	0.9900	C21—H21	0.9500
C9—H9B	0.9900	C21—C22	1.393 (2)
C9—C10	1.5162 (19)	C22—H22	0.9500
C10—C11	1.393 (2)		
C1—N1—C2	123.48 (12)	C15—C10—C11	118.84 (14)
C1—N1—H1	121.3 (18)	C10—C11—H11	119.7
C2—N1—H1	114.6 (18)	C12—C11—C10	120.66 (15)
C1—N2—C9	121.00 (12)	C12—C11—H11	119.7
C1—N2—C16	122.75 (13)	C11—C12—H12	120.0
C9—N2—C16	116.25 (11)	C13—C12—C11	120.03 (15)
N1—C1—S1	120.99 (11)	C13—C12—H12	120.0
N1—C1—N2	116.81 (13)	C12—C13—H13	120.2
N2—C1—S1	122.13 (11)	C14—C13—C12	119.64 (14)
N1—C2—H2A	109.2	C14—C13—H13	120.2
N1—C2—H2B	109.2	C13—C14—H14	119.8
N1—C2—C3	112.20 (11)	C13—C14—C15	120.39 (15)
H2A—C2—H2B	107.9	C15—C14—H14	119.8
C3—C2—H2A	109.2	C10—C15—H15	119.8
C3—C2—H2B	109.2	C14—C15—C10	120.43 (14)
C4—C3—C2	119.63 (14)	C14—C15—H15	119.8
C4—C3—C8	119.56 (15)	N2—C16—H16A	108.6
C8—C3—C2	120.79 (14)	N2—C16—H16B	108.6
C3—C4—H4	119.8	N2—C16—C17	114.82 (12)
C3—C4—C5	120.44 (18)	H16A—C16—H16B	107.5
C5—C4—H4	119.8	C17—C16—H16A	108.6
C4—C5—H5	120.1	C17—C16—H16B	108.6
C6—C5—C4	119.70 (19)	C18—C17—C16	118.44 (13)
C6—C5—H5	120.1	C22—C17—C16	122.47 (13)
C5—C6—H6	119.9	C22—C17—C18	119.05 (14)
C7—C6—C5	120.19 (17)	C17—C18—H18	119.7
C7—C6—H6	119.9	C19—C18—C17	120.51 (15)
C6—C7—H7	120.0	C19—C18—H18	119.7
C6—C7—C8	120.03 (18)	C18—C19—H19	119.9
C8—C7—H7	120.0	C18—C19—C20	120.19 (14)

C3—C8—C7	120.05 (17)	C20—C19—H19	119.9
C3—C8—H8	120.0	C19—C20—H20	120.3
C7—C8—H8	120.0	C21—C20—C19	119.47 (14)
N2—C9—H9A	108.7	C21—C20—H20	120.3
N2—C9—H9B	108.7	C20—C21—H21	119.8
N2—C9—C10	114.29 (12)	C20—C21—C22	120.48 (15)
H9A—C9—H9B	107.6	C22—C21—H21	119.8
C10—C9—H9A	108.7	C17—C22—C21	120.29 (14)
C10—C9—H9B	108.7	C17—C22—H22	119.9
C11—C10—C9	118.69 (13)	C21—C22—H22	119.9
C15—C10—C9	122.46 (13)		
N1—C2—C3—C4	57.70 (19)	C9—N2—C16—C17	-92.33 (15)
N1—C2—C3—C8	-123.79 (15)	C9—C10—C11—C12	-179.93 (13)
N2—C9—C10—C11	-179.60 (12)	C9—C10—C15—C14	178.87 (13)
N2—C9—C10—C15	1.06 (19)	C10—C11—C12—C13	0.9 (2)
N2—C16—C17—C18	164.26 (13)	C11—C10—C15—C14	-0.5 (2)
N2—C16—C17—C22	-18.1 (2)	C11—C12—C13—C14	-0.1 (2)
C1—N1—C2—C3	77.11 (17)	C12—C13—C14—C15	-0.9 (2)
C1—N2—C9—C10	95.30 (15)	C13—C14—C15—C10	1.2 (2)
C1—N2—C16—C17	88.11 (17)	C15—C10—C11—C12	-0.6 (2)
C2—N1—C1—S1	11.18 (19)	C16—N2—C1—S1	168.53 (10)
C2—N1—C1—N2	-171.82 (12)	C16—N2—C1—N1	-8.43 (19)
C2—C3—C4—C5	-179.84 (17)	C16—N2—C9—C10	-84.26 (14)
C2—C3—C8—C7	-179.52 (15)	C16—C17—C18—C19	178.27 (14)
C3—C4—C5—C6	-0.9 (3)	C16—C17—C22—C21	-177.37 (14)
C4—C3—C8—C7	-1.0 (2)	C17—C18—C19—C20	-0.6 (2)
C4—C5—C6—C7	-0.5 (3)	C18—C17—C22—C21	0.3 (2)
C5—C6—C7—C8	1.1 (3)	C18—C19—C20—C21	0.0 (2)
C6—C7—C8—C3	-0.4 (3)	C19—C20—C21—C22	0.8 (2)
C8—C3—C4—C5	1.6 (3)	C20—C21—C22—C17	-1.0 (2)
C9—N2—C1—S1	-11.01 (18)	C22—C17—C18—C19	0.5 (2)
C9—N2—C1—N1	172.03 (12)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1...S1 ⁱ	0.86 (3)	2.47 (3)	3.2044 (13)	145 (3)

Symmetry code: (i) *x*, -*y*+1/2, *z*+1/2.*N*-methyl-*N,N'*-Diphenylthiourea (2)

Crystal data

C₁₄H₁₄N₂S*M_r* = 242.33Triclinic, *P*1̄*a* = 9.8379 (6) Å*b* = 10.8014 (6) Å*c* = 13.2328 (6) Å α = 65.913 (5)° β = 87.752 (4)° γ = 84.059 (5)°*V* = 1276.82 (13) Å³*Z* = 4*F*(000) = 512

$D_x = 1.261 \text{ Mg m}^{-3}$
 Cu $K\alpha$ radiation, $\lambda = 1.54184 \text{ \AA}$
 Cell parameters from 5889 reflections
 $\theta = 3.6\text{--}73.7^\circ$

$\mu = 2.06 \text{ mm}^{-1}$
 $T = 100 \text{ K}$
 Block, clear colourless
 $0.26 \times 0.17 \times 0.13 \text{ mm}$

Data collection

SuperNova, Dual, Cu at home/near, Atlas diffractometer
 Radiation source: micro-focus sealed X-ray tube, SuperNova (Cu) X-ray Source
 Mirror monochromator
 Detector resolution: $10.4839 \text{ pixels mm}^{-1}$
 ω scans
 Absorption correction: gaussian (CrysAlisPro; Rigaku OD, 2019)

$T_{\min} = 0.687$, $T_{\max} = 1.000$
 12208 measured reflections
 4818 independent reflections
 4298 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$
 $\theta_{\max} = 73.9^\circ$, $\theta_{\min} = 3.7^\circ$
 $h = -11 \rightarrow 11$
 $k = -13 \rightarrow 12$
 $l = -16 \rightarrow 16$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.142$
 $S = 1.06$
 4818 reflections
 315 parameters
 0 restraints
 Primary atom site location: dual

Hydrogen site location: mixed
 H atoms treated by a mixture of independent and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0963P)^2 + 0.1879P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.58 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.25 \text{ e \AA}^{-3}$

Special details

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

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.87530 (5)	0.36219 (4)	-0.10166 (3)	0.03871 (15)
N1	0.74118 (16)	0.27410 (15)	0.09073 (12)	0.0352 (3)
N2	0.78346 (16)	0.49902 (15)	0.01836 (12)	0.0385 (3)
C1	0.79648 (17)	0.37896 (17)	0.00793 (13)	0.0344 (4)
C2	0.74183 (18)	0.13995 (17)	0.09466 (12)	0.0339 (4)
C3	0.61818 (19)	0.08717 (18)	0.09855 (13)	0.0370 (4)
H3	0.534808	0.141270	0.096050	0.044*
C4	0.6160 (2)	-0.04473 (19)	0.10612 (14)	0.0415 (4)
H4	0.531343	-0.081517	0.110231	0.050*
C5	0.7377 (2)	-0.12244 (19)	0.10765 (14)	0.0431 (4)
H5	0.736363	-0.211941	0.111191	0.052*
C6	0.8613 (2)	-0.06994 (19)	0.10404 (14)	0.0419 (4)
H6	0.944550	-0.123412	0.104909	0.050*
C7	0.86365 (19)	0.06084 (18)	0.09914 (13)	0.0383 (4)
H7	0.948411	0.095870	0.098860	0.046*
C8	0.8263 (2)	0.62333 (19)	-0.06957 (16)	0.0466 (4)

H8A	0.755527	0.661994	-0.127069	0.070*
H8B	0.912030	0.601735	-0.101872	0.070*
H8C	0.840149	0.689582	-0.038711	0.070*
C9	0.71401 (18)	0.51678 (17)	0.11024 (14)	0.0357 (4)
C10	0.78585 (18)	0.49184 (18)	0.20553 (15)	0.0379 (4)
H10	0.879437	0.457338	0.212657	0.045*
C11	0.7198 (2)	0.51775 (19)	0.29091 (15)	0.0429 (4)
H11	0.768623	0.501230	0.356392	0.052*
C12	0.5839 (2)	0.5672 (2)	0.28090 (17)	0.0461 (4)
H12	0.539256	0.584186	0.339560	0.055*
C13	0.5120 (2)	0.5923 (2)	0.18474 (19)	0.0470 (4)
H13	0.418349	0.626521	0.177738	0.056*
C14	0.5773 (2)	0.56728 (19)	0.09959 (17)	0.0420 (4)
H14	0.528599	0.584619	0.033840	0.050*
S21	0.52714 (5)	0.19752 (5)	0.30490 (4)	0.04442 (16)
N21	0.47106 (17)	0.1816 (2)	0.50962 (13)	0.0464 (4)
N22	0.69539 (17)	0.14244 (18)	0.47379 (14)	0.0456 (4)
C21	0.5669 (2)	0.17492 (19)	0.43396 (15)	0.0408 (4)
C22	0.33316 (19)	0.23455 (19)	0.49771 (14)	0.0380 (4)
C23	0.2535 (2)	0.1894 (2)	0.59315 (14)	0.0438 (4)
H23	0.292688	0.122901	0.660265	0.053*
C24	0.1187 (2)	0.2401 (2)	0.59142 (16)	0.0491 (5)
H24	0.065365	0.208041	0.656896	0.059*
C25	0.0610 (2)	0.3379 (2)	0.49387 (18)	0.0520 (5)
H25	-0.031833	0.373337	0.492102	0.062*
C26	0.1397 (2)	0.3832 (2)	0.39947 (17)	0.0503 (5)
H26	0.099883	0.449752	0.332619	0.060*
C27	0.2752 (2)	0.3338 (2)	0.39998 (15)	0.0431 (4)
H27	0.328299	0.367067	0.334463	0.052*
C28	0.8129 (2)	0.1216 (2)	0.4083 (2)	0.0526 (5)
H28A	0.862359	0.203300	0.379216	0.079*
H28B	0.780515	0.104245	0.346634	0.079*
H28C	0.874188	0.043352	0.455457	0.079*
C29	0.72661 (18)	0.1336 (2)	0.58213 (15)	0.0418 (4)
C30	0.7548 (2)	0.0084 (2)	0.6690 (2)	0.0541 (5)
H30	0.756557	-0.072996	0.657398	0.065*
C31	0.7807 (3)	0.0024 (2)	0.7737 (2)	0.0621 (6)
H31	0.798209	-0.083740	0.833900	0.074*
H1	0.688 (3)	0.290 (3)	0.138 (3)	0.074*
H21	0.494 (3)	0.135 (3)	0.580 (3)	0.074*
C32	0.7813 (2)	0.1195 (2)	0.79115 (17)	0.0497 (5)
H32	0.798612	0.114241	0.862995	0.060*
C33	0.7567 (2)	0.2446 (2)	0.70368 (16)	0.0438 (4)
H33	0.759655	0.325717	0.714838	0.053*
C34	0.72752 (19)	0.2523 (2)	0.59942 (15)	0.0430 (4)
H34	0.708181	0.338601	0.539811	0.052*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0441 (3)	0.0393 (3)	0.0306 (2)	-0.00590 (17)	0.00684 (17)	-0.01224 (18)
N1	0.0400 (8)	0.0323 (7)	0.0335 (7)	-0.0053 (5)	0.0067 (6)	-0.0139 (6)
N2	0.0462 (8)	0.0326 (7)	0.0349 (7)	-0.0076 (6)	0.0072 (6)	-0.0116 (6)
C1	0.0316 (8)	0.0362 (9)	0.0331 (7)	-0.0041 (6)	0.0001 (6)	-0.0117 (6)
C2	0.0425 (9)	0.0322 (8)	0.0262 (7)	-0.0054 (6)	0.0031 (6)	-0.0109 (6)
C3	0.0413 (9)	0.0372 (9)	0.0303 (7)	-0.0037 (7)	-0.0002 (6)	-0.0115 (6)
C4	0.0521 (11)	0.0398 (9)	0.0326 (8)	-0.0118 (8)	-0.0010 (7)	-0.0128 (7)
C5	0.0642 (12)	0.0328 (9)	0.0318 (8)	-0.0049 (8)	0.0016 (8)	-0.0128 (7)
C6	0.0506 (11)	0.0384 (9)	0.0323 (8)	0.0028 (7)	0.0058 (7)	-0.0118 (7)
C7	0.0419 (9)	0.0381 (9)	0.0314 (8)	-0.0045 (7)	0.0036 (7)	-0.0108 (7)
C8	0.0623 (12)	0.0345 (9)	0.0402 (9)	-0.0127 (8)	0.0105 (8)	-0.0116 (7)
C9	0.0397 (9)	0.0299 (8)	0.0378 (8)	-0.0068 (6)	0.0046 (7)	-0.0138 (7)
C10	0.0365 (9)	0.0344 (9)	0.0385 (8)	-0.0038 (6)	0.0026 (7)	-0.0108 (7)
C11	0.0516 (11)	0.0408 (10)	0.0353 (8)	-0.0099 (8)	0.0040 (7)	-0.0133 (7)
C12	0.0507 (11)	0.0430 (10)	0.0485 (10)	-0.0098 (8)	0.0144 (8)	-0.0225 (8)
C13	0.0361 (10)	0.0461 (11)	0.0633 (12)	-0.0038 (7)	0.0051 (8)	-0.0273 (9)
C14	0.0402 (10)	0.0383 (9)	0.0501 (10)	-0.0039 (7)	-0.0037 (8)	-0.0203 (8)
S21	0.0442 (3)	0.0584 (3)	0.0376 (2)	-0.0158 (2)	0.00781 (19)	-0.0247 (2)
N21	0.0393 (9)	0.0629 (11)	0.0288 (7)	0.0000 (7)	0.0020 (6)	-0.0115 (7)
N22	0.0390 (9)	0.0546 (10)	0.0426 (8)	-0.0017 (7)	0.0057 (6)	-0.0203 (7)
C21	0.0415 (10)	0.0406 (9)	0.0395 (9)	-0.0068 (7)	0.0062 (7)	-0.0154 (7)
C22	0.0373 (9)	0.0443 (9)	0.0324 (8)	-0.0068 (7)	0.0013 (7)	-0.0150 (7)
C23	0.0438 (10)	0.0516 (11)	0.0301 (8)	-0.0021 (8)	0.0011 (7)	-0.0112 (7)
C24	0.0422 (10)	0.0597 (12)	0.0366 (9)	-0.0036 (8)	0.0076 (7)	-0.0116 (8)
C25	0.0364 (10)	0.0549 (12)	0.0515 (11)	-0.0008 (8)	0.0011 (8)	-0.0092 (9)
C26	0.0444 (11)	0.0491 (11)	0.0417 (9)	-0.0040 (8)	-0.0022 (8)	-0.0025 (8)
C27	0.0417 (10)	0.0465 (10)	0.0340 (8)	-0.0088 (7)	0.0034 (7)	-0.0084 (7)
C28	0.0433 (11)	0.0587 (12)	0.0612 (12)	-0.0032 (9)	0.0114 (9)	-0.0312 (10)
C29	0.0321 (9)	0.0482 (10)	0.0412 (9)	-0.0036 (7)	0.0010 (7)	-0.0143 (8)
C30	0.0559 (13)	0.0411 (11)	0.0601 (12)	-0.0092 (9)	-0.0146 (10)	-0.0131 (9)
C31	0.0719 (15)	0.0467 (12)	0.0534 (12)	-0.0157 (10)	-0.0225 (11)	-0.0017 (9)
C32	0.0472 (11)	0.0539 (12)	0.0407 (9)	-0.0128 (8)	-0.0050 (8)	-0.0094 (8)
C33	0.0395 (10)	0.0489 (11)	0.0407 (9)	-0.0052 (7)	0.0061 (7)	-0.0161 (8)
C34	0.0399 (9)	0.0457 (10)	0.0356 (8)	0.0006 (7)	0.0055 (7)	-0.0101 (7)

Geometric parameters (\AA , $^\circ$)

S1—C1	1.6835 (17)	S21—C21	1.6798 (19)
N1—C1	1.359 (2)	N21—C21	1.367 (2)
N1—C2	1.428 (2)	N21—C22	1.405 (2)
N1—H1	0.86 (4)	N21—H21	0.88 (3)
N2—C1	1.352 (2)	N22—C21	1.345 (3)
N2—C8	1.462 (2)	N22—C28	1.472 (2)
N2—C9	1.442 (2)	N22—C29	1.441 (3)
C2—C3	1.386 (3)	C22—C23	1.395 (3)

C2—C7	1.386 (2)	C22—C27	1.395 (3)
C3—H3	0.9500	C23—H23	0.9500
C3—C4	1.389 (3)	C23—C24	1.379 (3)
C4—H4	0.9500	C24—H24	0.9500
C4—C5	1.385 (3)	C24—C25	1.387 (3)
C5—H5	0.9500	C25—H25	0.9500
C5—C6	1.385 (3)	C25—C26	1.379 (3)
C6—H6	0.9500	C26—H26	0.9500
C6—C7	1.390 (3)	C26—C27	1.383 (3)
C7—H7	0.9500	C27—H27	0.9500
C8—H8A	0.9800	C28—H28A	0.9800
C8—H8B	0.9800	C28—H28B	0.9800
C8—H8C	0.9800	C28—H28C	0.9800
C9—C10	1.384 (3)	C29—C30	1.381 (3)
C9—C14	1.388 (3)	C29—C34	1.392 (3)
C10—H10	0.9500	C30—H30	0.9500
C10—C11	1.393 (3)	C30—C31	1.393 (4)
C11—H11	0.9500	C31—H31	0.9500
C11—C12	1.377 (3)	C31—C32	1.376 (4)
C12—H12	0.9500	C32—H32	0.9500
C12—C13	1.394 (3)	C32—C33	1.379 (3)
C13—H13	0.9500	C33—H33	0.9500
C13—C14	1.382 (3)	C33—C34	1.387 (3)
C14—H14	0.9500	C34—H34	0.9500
C1—N1—C2	124.62 (14)	C21—N21—C22	131.73 (16)
C1—N1—H1	120 (2)	C21—N21—H21	116 (2)
C2—N1—H1	114 (2)	C22—N21—H21	112 (2)
C1—N2—C8	121.52 (15)	C21—N22—C28	122.70 (17)
C1—N2—C9	122.74 (14)	C21—N22—C29	121.26 (16)
C9—N2—C8	115.42 (14)	C29—N22—C28	115.95 (17)
N1—C1—S1	122.60 (13)	N21—C21—S21	123.08 (15)
N2—C1—S1	121.65 (13)	N22—C21—S21	122.99 (14)
N2—C1—N1	115.75 (15)	N22—C21—N21	113.87 (17)
C3—C2—N1	118.94 (15)	C23—C22—N21	116.39 (16)
C3—C2—C7	120.07 (17)	C23—C22—C27	119.09 (17)
C7—C2—N1	120.94 (16)	C27—C22—N21	124.39 (17)
C2—C3—H3	120.0	C22—C23—H23	119.5
C2—C3—C4	120.09 (17)	C24—C23—C22	120.92 (17)
C4—C3—H3	120.0	C24—C23—H23	119.5
C3—C4—H4	120.1	C23—C24—H24	120.1
C5—C4—C3	119.83 (19)	C23—C24—C25	119.87 (18)
C5—C4—H4	120.1	C25—C24—H24	120.1
C4—C5—H5	119.9	C24—C25—H25	120.3
C4—C5—C6	120.12 (18)	C26—C25—C24	119.35 (19)
C6—C5—H5	119.9	C26—C25—H25	120.3
C5—C6—H6	119.9	C25—C26—H26	119.3
C5—C6—C7	120.13 (17)	C25—C26—C27	121.47 (18)

C7—C6—H6	119.9	C27—C26—H26	119.3
C2—C7—C6	119.73 (18)	C22—C27—H27	120.3
C2—C7—H7	120.1	C26—C27—C22	119.30 (17)
C6—C7—H7	120.1	C26—C27—H27	120.3
N2—C8—H8A	109.5	N22—C28—H28A	109.5
N2—C8—H8B	109.5	N22—C28—H28B	109.5
N2—C8—H8C	109.5	N22—C28—H28C	109.5
H8A—C8—H8B	109.5	H28A—C28—H28B	109.5
H8A—C8—H8C	109.5	H28A—C28—H28C	109.5
H8B—C8—H8C	109.5	H28B—C28—H28C	109.5
C10—C9—N2	119.96 (16)	C30—C29—N22	120.46 (19)
C10—C9—C14	120.40 (16)	C30—C29—C34	119.83 (19)
C14—C9—N2	119.52 (16)	C34—C29—N22	119.71 (17)
C9—C10—H10	120.3	C29—C30—H30	120.3
C9—C10—C11	119.43 (17)	C29—C30—C31	119.4 (2)
C11—C10—H10	120.3	C31—C30—H30	120.3
C10—C11—H11	119.8	C30—C31—H31	119.5
C12—C11—C10	120.35 (18)	C32—C31—C30	120.9 (2)
C12—C11—H11	119.8	C32—C31—H31	119.5
C11—C12—H12	120.0	C31—C32—H32	120.2
C11—C12—C13	120.04 (17)	C31—C32—C33	119.6 (2)
C13—C12—H12	120.0	C33—C32—H32	120.2
C12—C13—H13	120.1	C32—C33—H33	119.9
C14—C13—C12	119.84 (18)	C32—C33—C34	120.2 (2)
C14—C13—H13	120.1	C34—C33—H33	119.9
C9—C14—H14	120.0	C29—C34—H34	120.0
C13—C14—C9	119.94 (18)	C33—C34—C29	120.03 (18)
C13—C14—H14	120.0	C33—C34—H34	120.0
N1—C2—C3—C4	-177.86 (15)	N21—C22—C23—C24	176.9 (2)
N1—C2—C7—C6	179.34 (15)	N21—C22—C27—C26	-176.8 (2)
N2—C9—C10—C11	175.88 (16)	N22—C29—C30—C31	-178.1 (2)
N2—C9—C14—C13	-176.16 (17)	N22—C29—C34—C33	179.60 (17)
C1—N1—C2—C3	-120.49 (18)	C21—N21—C22—C23	161.2 (2)
C1—N1—C2—C7	62.2 (2)	C21—N21—C22—C27	-23.1 (4)
C1—N2—C9—C10	89.0 (2)	C21—N22—C29—C30	105.4 (2)
C1—N2—C9—C14	-95.1 (2)	C21—N22—C29—C34	-74.2 (3)
C2—N1—C1—S1	-0.6 (2)	C22—N21—C21—S21	-16.0 (3)
C2—N1—C1—N2	178.92 (16)	C22—N21—C21—N22	166.9 (2)
C2—C3—C4—C5	-1.2 (3)	C22—C23—C24—C25	-0.4 (3)
C3—C2—C7—C6	2.0 (2)	C23—C22—C27—C26	-1.2 (3)
C3—C4—C5—C6	1.4 (3)	C23—C24—C25—C26	0.1 (4)
C4—C5—C6—C7	0.2 (3)	C24—C25—C26—C27	-0.4 (4)
C5—C6—C7—C2	-1.9 (3)	C25—C26—C27—C22	0.9 (3)
C7—C2—C3—C4	-0.5 (2)	C27—C22—C23—C24	1.0 (3)
C8—N2—C1—S1	5.5 (2)	C28—N22—C21—S21	-1.0 (3)
C8—N2—C1—N1	-174.01 (17)	C28—N22—C21—N21	176.15 (19)
C8—N2—C9—C10	-97.4 (2)	C28—N22—C29—C30	-78.0 (3)

C8—N2—C9—C14	78.5 (2)	C28—N22—C29—C34	102.4 (2)
C9—N2—C1—S1	178.71 (13)	C29—N22—C21—S21	175.44 (15)
C9—N2—C1—N1	-0.8 (2)	C29—N22—C21—N21	-7.4 (3)
C9—C10—C11—C12	0.3 (3)	C29—C30—C31—C32	-1.4 (4)
C10—C9—C14—C13	-0.3 (3)	C30—C29—C34—C33	0.0 (3)
C10—C11—C12—C13	-0.4 (3)	C30—C31—C32—C33	-0.3 (4)
C11—C12—C13—C14	0.1 (3)	C31—C32—C33—C34	1.9 (3)
C12—C13—C14—C9	0.2 (3)	C32—C33—C34—C29	-1.7 (3)
C14—C9—C10—C11	0.0 (3)	C34—C29—C30—C31	1.5 (3)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
N1—H1...S21	0.86 (3)	2.58 (3)	3.3360 (16)	148 (3)

N,N-Di-*n*-butyl-*N'*-phenylthiourea (3)*Crystal data*C₁₅H₂₄N₂S*M_r* = 264.42Trigonal, *R*3̄*a* = 25.5231 (3) Å*c* = 12.6225 (2) Å*V* = 7121.0 (2) Å³*Z* = 18*F*(000) = 2592*D_x* = 1.110 Mg m⁻³Cu *Kα* radiation, λ = 1.54184 Å

Cell parameters from 8527 reflections

θ = 3.4–73.3°

μ = 1.69 mm⁻¹*T* = 100 K

Block, clear colourless

0.42 × 0.26 × 0.18 mm

Data collection

SuperNova, Dual, Cu at home/near, Atlas diffractometer

Radiation source: micro-focus sealed X-ray tube, SuperNova (Cu) X-ray Source

Mirror monochromator

Detector resolution: 10.4839 pixels mm⁻¹

ω scans

Absorption correction: gaussian (CrysAlisPro; Rigaku OD, 2019)

T_{min} = 0.479, *T_{max}* = 1.000

14921 measured reflections

3151 independent reflections

3004 reflections with *I* > 2σ(*I*)*R_{int}* = 0.021θ_{max} = 73.7°, θ_{min} = 3.5°*h* = -31→31*k* = -28→28*l* = -15→15*Refinement*Refinement on *F*²

Least-squares matrix: full

R[*F*² > 2σ(*F*²)] = 0.032*wR*(*F*²) = 0.087*S* = 1.06

3151 reflections

168 parameters

0 restraints

Primary atom site location: dual

Hydrogen site location: mixed

H atoms treated by a mixture of independent and constrained refinement

w = 1/[σ²(*F_o*²) + (0.0479*P*)² + 4.6459*P*]where *P* = (*F_o*² + 2*F_c*²)/3(Δ/σ)_{max} = 0.001Δρ_{max} = 0.25 e Å⁻³Δρ_{min} = -0.18 e Å⁻³

Special details

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

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.52061 (2)	0.89087 (2)	0.56407 (2)	0.03089 (10)
N1	0.54435 (4)	0.80126 (4)	0.59489 (8)	0.0307 (2)
N2	0.60326 (4)	0.88769 (4)	0.69020 (7)	0.0277 (2)
C1	0.55829 (5)	0.85861 (5)	0.61987 (8)	0.0269 (2)
C2	0.50126 (5)	0.76273 (5)	0.51835 (9)	0.0300 (2)
C3	0.45844 (5)	0.70441 (6)	0.54908 (10)	0.0346 (3)
H3	0.456357	0.692296	0.620843	0.041*
C4	0.41868 (6)	0.66385 (6)	0.47470 (12)	0.0409 (3)
H4	0.389778	0.623828	0.495424	0.049*
C5	0.42119 (6)	0.68178 (6)	0.37040 (11)	0.0415 (3)
H5	0.393836	0.654162	0.319656	0.050*
C6	0.46357 (6)	0.73993 (7)	0.34018 (10)	0.0404 (3)
H6	0.464950	0.752227	0.268704	0.049*
C7	0.50412 (5)	0.78049 (6)	0.41355 (10)	0.0350 (3)
H7	0.533631	0.820148	0.392150	0.042*
C8	0.61573 (5)	0.94484 (5)	0.73935 (9)	0.0319 (2)
H8A	0.602928	0.966811	0.690774	0.038*
H8B	0.659778	0.970459	0.751515	0.038*
C9	0.58233 (5)	0.93370 (5)	0.84445 (9)	0.0333 (3)
H9A	0.538255	0.909261	0.831385	0.040*
H9B	0.593836	0.910012	0.891379	0.040*
C10	0.59590 (7)	0.99172 (6)	0.90032 (10)	0.0426 (3)
H10A	0.582272	1.014382	0.855269	0.051*
H10B	0.640170	1.017203	0.910118	0.051*
C11	0.56494 (7)	0.97991 (7)	1.00770 (12)	0.0493 (3)
H11A	0.520992	0.956005	0.998160	0.074*
H11B	0.575502	1.018545	1.041429	0.074*
H11C	0.578367	0.957572	1.052685	0.074*
H1	0.5540 (9)	0.7814 (9)	0.6385 (15)	0.059*
C12	0.64531 (5)	0.86653 (5)	0.71978 (9)	0.0288 (2)
H12A	0.622637	0.821985	0.727847	0.035*
H12B	0.664311	0.884432	0.788746	0.035*
C13	0.69452 (5)	0.88425 (6)	0.63577 (9)	0.0342 (3)
H13A	0.675206	0.869399	0.565789	0.041*
H13B	0.719121	0.928910	0.632074	0.041*
C14	0.73574 (6)	0.85867 (6)	0.65902 (10)	0.0359 (3)
H14A	0.711309	0.813967	0.661602	0.043*
H14B	0.754711	0.873062	0.729373	0.043*
C15	0.78511 (6)	0.87735 (7)	0.57560 (12)	0.0477 (3)

H15A	0.766520	0.863934	0.505610	0.071*
H15B	0.809652	0.858661	0.591775	0.071*
H15C	0.810909	0.921480	0.575771	0.071*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.03175 (16)	0.03319 (16)	0.03491 (17)	0.02161 (12)	0.00062 (10)	0.00343 (10)
N1	0.0290 (5)	0.0276 (5)	0.0384 (5)	0.0163 (4)	-0.0062 (4)	-0.0022 (4)
N2	0.0267 (4)	0.0242 (4)	0.0329 (5)	0.0133 (4)	-0.0008 (4)	0.0005 (4)
C1	0.0244 (5)	0.0266 (5)	0.0307 (5)	0.0136 (4)	0.0036 (4)	0.0030 (4)
C2	0.0241 (5)	0.0301 (6)	0.0394 (6)	0.0164 (5)	-0.0024 (4)	-0.0056 (4)
C3	0.0294 (6)	0.0314 (6)	0.0453 (6)	0.0170 (5)	-0.0027 (5)	-0.0012 (5)
C4	0.0296 (6)	0.0315 (6)	0.0594 (8)	0.0137 (5)	-0.0045 (5)	-0.0067 (5)
C5	0.0308 (6)	0.0444 (7)	0.0491 (7)	0.0186 (6)	-0.0067 (5)	-0.0172 (6)
C6	0.0331 (6)	0.0525 (8)	0.0360 (6)	0.0217 (6)	0.0004 (5)	-0.0085 (5)
C7	0.0270 (5)	0.0380 (6)	0.0384 (6)	0.0150 (5)	0.0030 (4)	-0.0026 (5)
C8	0.0339 (6)	0.0232 (5)	0.0374 (6)	0.0134 (5)	-0.0031 (5)	-0.0009 (4)
C9	0.0316 (6)	0.0290 (6)	0.0391 (6)	0.0151 (5)	-0.0025 (5)	-0.0028 (5)
C10	0.0620 (9)	0.0358 (7)	0.0368 (6)	0.0294 (6)	-0.0076 (6)	-0.0041 (5)
C11	0.0566 (9)	0.0548 (8)	0.0453 (7)	0.0344 (7)	-0.0024 (6)	-0.0111 (6)
C12	0.0263 (5)	0.0261 (5)	0.0338 (5)	0.0130 (4)	-0.0032 (4)	0.0016 (4)
C13	0.0315 (6)	0.0361 (6)	0.0365 (6)	0.0180 (5)	0.0006 (5)	0.0029 (5)
C14	0.0347 (6)	0.0396 (6)	0.0372 (6)	0.0214 (5)	-0.0042 (5)	-0.0046 (5)
C15	0.0337 (7)	0.0558 (8)	0.0531 (8)	0.0220 (6)	0.0010 (6)	-0.0082 (6)

Geometric parameters (Å, °)

S1—C1	1.7004 (11)	C9—H9A	0.9900
N1—C1	1.3594 (15)	C9—H9B	0.9900
N1—C2	1.4241 (15)	C9—C10	1.5157 (17)
N1—H1	0.86 (2)	C10—H10A	0.9900
N2—C1	1.3432 (15)	C10—H10B	0.9900
N2—C8	1.4663 (14)	C10—C11	1.521 (2)
N2—C12	1.4705 (14)	C11—H11A	0.9800
C2—C3	1.3907 (17)	C11—H11B	0.9800
C2—C7	1.3885 (17)	C11—H11C	0.9800
C3—H3	0.9500	C12—H12A	0.9900
C3—C4	1.3901 (18)	C12—H12B	0.9900
C4—H4	0.9500	C12—C13	1.5293 (16)
C4—C5	1.385 (2)	C13—H13A	0.9900
C5—H5	0.9500	C13—H13B	0.9900
C5—C6	1.383 (2)	C13—C14	1.5186 (17)
C6—H6	0.9500	C14—H14A	0.9900
C6—C7	1.3888 (18)	C14—H14B	0.9900
C7—H7	0.9500	C14—C15	1.5242 (18)
C8—H8A	0.9900	C15—H15A	0.9800
C8—H8B	0.9900	C15—H15B	0.9800

C8—C9	1.5249 (17)	C15—H15C	0.9800
C1—N1—C2	126.66 (10)	C10—C9—H9B	109.0
C1—N1—H1	119.1 (13)	C9—C10—H10A	109.2
C2—N1—H1	112.1 (13)	C9—C10—H10B	109.2
C1—N2—C8	121.99 (9)	C9—C10—C11	112.25 (12)
C1—N2—C12	122.95 (9)	H10A—C10—H10B	107.9
C8—N2—C12	115.00 (9)	C11—C10—H10A	109.2
N1—C1—S1	121.22 (8)	C11—C10—H10B	109.2
N2—C1—S1	122.70 (8)	C10—C11—H11A	109.5
N2—C1—N1	116.08 (10)	C10—C11—H11B	109.5
C3—C2—N1	118.18 (11)	C10—C11—H11C	109.5
C7—C2—N1	121.64 (11)	H11A—C11—H11B	109.5
C7—C2—C3	120.00 (11)	H11A—C11—H11C	109.5
C2—C3—H3	120.0	H11B—C11—H11C	109.5
C4—C3—C2	119.94 (12)	N2—C12—H12A	109.4
C4—C3—H3	120.0	N2—C12—H12B	109.4
C3—C4—H4	120.0	N2—C12—C13	110.96 (9)
C5—C4—C3	120.01 (12)	H12A—C12—H12B	108.0
C5—C4—H4	120.0	C13—C12—H12A	109.4
C4—C5—H5	120.0	C13—C12—H12B	109.4
C6—C5—C4	119.94 (12)	C12—C13—H13A	109.1
C6—C5—H5	120.0	C12—C13—H13B	109.1
C5—C6—H6	119.8	H13A—C13—H13B	107.8
C5—C6—C7	120.48 (13)	C14—C13—C12	112.46 (10)
C7—C6—H6	119.8	C14—C13—H13A	109.1
C2—C7—C6	119.61 (12)	C14—C13—H13B	109.1
C2—C7—H7	120.2	C13—C14—H14A	109.2
C6—C7—H7	120.2	C13—C14—H14B	109.2
N2—C8—H8A	109.4	C13—C14—C15	111.96 (11)
N2—C8—H8B	109.4	H14A—C14—H14B	107.9
N2—C8—C9	111.07 (9)	C15—C14—H14A	109.2
H8A—C8—H8B	108.0	C15—C14—H14B	109.2
C9—C8—H8A	109.4	C14—C15—H15A	109.5
C9—C8—H8B	109.4	C14—C15—H15B	109.5
C8—C9—H9A	109.0	C14—C15—H15C	109.5
C8—C9—H9B	109.0	H15A—C15—H15B	109.5
H9A—C9—H9B	107.8	H15A—C15—H15C	109.5
C10—C9—C8	112.91 (10)	H15B—C15—H15C	109.5
C10—C9—H9A	109.0		
N1—C2—C3—C4	-174.90 (10)	C3—C4—C5—C6	0.45 (19)
N1—C2—C7—C6	175.74 (11)	C4—C5—C6—C7	0.57 (19)
N2—C8—C9—C10	177.68 (10)	C5—C6—C7—C2	-1.16 (19)
N2—C12—C13—C14	-175.23 (10)	C7—C2—C3—C4	0.27 (17)
C1—N1—C2—C3	-129.37 (12)	C8—N2—C1—S1	11.78 (15)
C1—N1—C2—C7	55.54 (16)	C8—N2—C1—N1	-168.43 (10)
C1—N2—C8—C9	92.75 (12)	C8—N2—C12—C13	-97.50 (11)

C1—N2—C12—C13	79.64 (13)	C8—C9—C10—C11	-176.92 (11)
C2—N1—C1—S1	3.80 (16)	C12—N2—C1—S1	-165.16 (8)
C2—N1—C1—N2	-176.00 (10)	C12—N2—C1—N1	14.63 (15)
C2—C3—C4—C5	-0.87 (18)	C12—N2—C8—C9	-90.08 (11)
C3—C2—C7—C6	0.74 (17)	C12—C13—C14—C15	-179.20 (11)

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N1—H1 \cdots S1 ⁱ	0.86 (2)	2.62 (2)	3.4656 (11)	167 (2)
C12—H12 <i>A</i> \cdots S1 ⁱ	0.99	2.67	3.6588 (13)	174

Symmetry code: (i) $y-1/3, -x+y+1/3, -z+4/3$.