



2-{3-Methyl-2-[(2Z)-pent-2-en-1-yl]cyclopent-2-en-1-ylidene}-N-phenylhydrazinecarbothioamide. Corrigendum

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In the paper by Oliveira *et al.* [*IUCrData* (2023), **8**, x230971], there was an error in the name of the first author.

The name of the first author in the paper by Oliveira *et al.* (2023) is incorrect and should be 'Adriano Bof de Oliveira', as given above.

References

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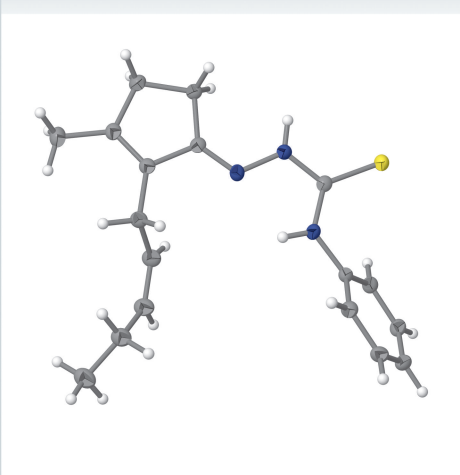
Keywords: *cis*-jasmane; 4-phenylthiosemicarbazone; thiosemicarbazone; jasmane; crystal structure; Hirshfeld analysis..

CCDC reference: 2304274

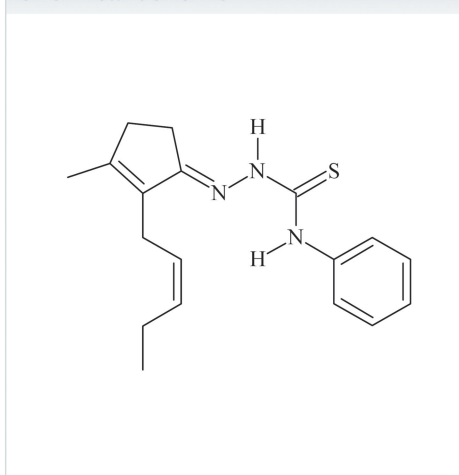
Structural data: full structural data are available from iucrdata.iucr.org

The hydrochloric acid-catalyzed equimolar reaction between *cis*-jasmane and 4-phenylthiosemicarbazide yielded the title compound, C₁₈H₂₃N₃S (common name: *cis*-jasmane 4-phenylthiosemicarbazone). Concerning the hydrogen bonding, an N—H···N intramolecular interaction is observed, forming a ring with graph-set motif *S*(5). In the crystal, the molecules are connected into centrosymmetric dimers by pairs of N—H···S and C—H···S interactions, forming rings of graph-set motifs *R*₂²(8) and *R*₂¹(7), with the sulfur atoms acting as double acceptors. The thiosemicarbazone entity is approximately planar, with the maximum deviation from the mean plane through the N/N/C/S/N atoms being 0.0376 (9) Å (the r.m.s.d. amounts to 0.0234 Å). The molecule is substantially twisted as indicated by the dihedral angle between the thiosemicarbazone fragment and the phenyl ring, which amounts to 56.1 (5)°, and because of the jasmane fragment, which bears a chain with *sp*³-hybridized carbon atoms in the structure. The Hirshfeld surface analysis indicates that the major contributions for the crystal cohesion are: H···H (65.3%), H···C/C···H (16.2%), H···S/S···H (10.9%) and H···N/N···H (5.5%).

3D view



Chemical scheme



Structure description

Thiosemicarbazone derivatives (**TSCs**), which are characterized by the [*R*₁*R*₂C=N—N(H)—C(=S)—NR₃R₄] functional group, were reported more than a century ago (Freund & Schander, 1902), while the synthesis of jasmane derivatives can be traced back to the early 1930s (Ruzicka & Pfeiffer, 1933). Concerning **TSC** chemistry, thiosemicarbazone molecules are the major product of the reaction between thiosemicarbazide derivatives [H₂N—N(H)—C(=S)—NR₃R₄] and aldehydes or ketones [*R*₁*R*₂C=O]. Thiosemicarbazides have been employed as analytical reagents in organic



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data reports

chemistry for the detection of the $[R_1R_2C=O]$ functional group by a condensation reaction through nucleophilic attack of the $[H_2N-]$ thiosemicarbazide fragment on the carbonyl group. Thiosemicarbazone chemistry gained new perspectives in the mid-1940s when some derivatives were pointed out in *in vitro* essays to be tuberculostatic agents (Domagk *et al.*, 1946). From these early times, this chemistry evolved into a large class of compounds with a wide range of applications across several scientific disciplines. The facile experimental procedure for the synthesis, combined with the vast structural diversity of the starting materials, *i.e.*, aldehydes and ketones, lead to a large number of **TSCs**. As a result of their molecular structure and the respective Lewis basicity (nitrogen atoms, with some more *hard* character, and the *soft* sulfur atom), allowing for chemical bonding with different metal centers in diverse modes, *e.g.*, bridging, chelating or terminal, thiosemicarbazones found several applications in coordination chemistry. For the synergetic effect of thiosemicarbazones and metal centers, see: Lobana *et al.* (2009). For the application on diagnostic medical imaging of **TSC** complexes, see: Dilworth & Hueting (2012) and for the application of **TSC** coordination compounds on theranostics, see: Parrilha *et al.* (2022). For electrocatalytic hydrogen production using a Pd^{II} complex with the 4-bis[4-(*p*-methoxyphenyl)thiosemicarbazone]]-2,3-butane derivative, which is relevant for the energy research today, see: Straistari *et al.* (2018). For biological applications of **TSCs** and their complexes, see: Singh *et al.* (2023). For the antifungal activity and the crystal structure of the non-substituted *cis*-jasmone thiosemicarbazone, see: Orsoni *et al.*, (2020) and for another report concerning the fungistatic effect of this **TSC** derivative, see: Jamiołkowska *et al.* (2022). For the application of thiosemicarbazones complexes as single-molecule precursors in the synthesis of nanostructured metal sulfides, see: Palve & Garje (2011) for ZnS, Pawar *et al.* (2016) for CdS and Pawar & Garje (2015) for CoS nanocrystalline materials. Regarding the use of a **TSC** on the formation of palladium nanoparticles for the Suzuki–Miyaura cross-coupling catalytic application, see: Kovala-Demertzi *et al.* (2008). Finally, to cite another example of their applications, thiosemicarbazones are employed as corrosion inhibitors. For an experimental and theoretical study regarding the corro-

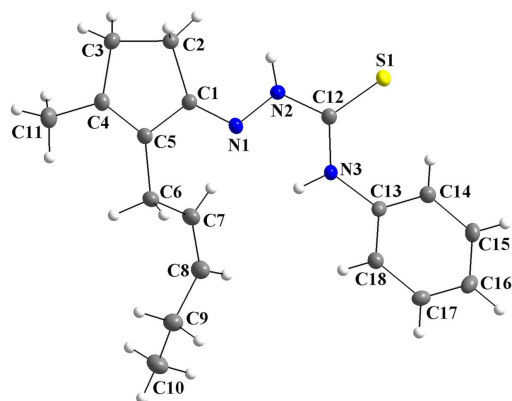


Figure 1
The molecular structure of the title compound, showing the atom labeling and displacement ellipsoids drawn at the 40% probability level.

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$C2-H2B\cdots S1^i$	0.97	2.96	3.4640 (16)	113
$N2-H2\cdots S1^i$	0.86	2.72	3.5757 (13)	177
$N3-H3\cdots N1$	0.86	2.11	2.5457 (18)	111

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

sion-inhibitory property of **TSCs** applied for carbon steel AISI 1020 in a hydrochloric acid medium, see: Goulart *et al.* (2013). For a theoretical approach of **TSC** dimers as corrosion inhibitors, see: Silva & Martínez-Huitle (2021).

As part of our interest in this chemistry, we report herein the synthesis, crystal structure and Hirshfeld analysis of the *cis*-jasmone 4-phenylthiosemicarbazone.

For the title compound, the molecular structure matches the asymmetric unit, with all atoms being located in general positions (Fig. 1). The thiosemicarbazone fragment is almost planar, with the maximum deviation from the mean plane through the N1/N2/C12/S1/N3 group being 0.0376 (9) Å for N2 and the r.m.s.d. for the selected atoms amounting to 0.0234 Å. The torsion angles of the N1–N2–C12–S1 and N1–N2–C12–N3 chains amount to 176.3 (1) and -5.2 (2)°. The C1–C5 pentagonal ring is almost planar, as the maximum deviation from the mean plane through the carbon atoms is 0.0117 (1) Å for C5 and the respective r.m.s.d. amounts to 0.0080 Å. The molecule is not planar because of the dihedral angle between the thiosemicarbazone entity and the phenyl ring, which is 56.1 (5)°, and due to the sp^3 -hybridized carbon atoms, *e.g.*, C6 and C9 in the jasmone fragment. In addition, an N3–H3···N1 intramolecular hydrogen bond is observed (Fig. 2, Table 1), with graph-set motif $S(5)$, which contributes to stabilize the molecular structure.

In the crystal, the molecules are connected into centrosymmetric dimers by pairs of N2–H2···S1ⁱ interactions, which form rings of graph-set motif $R_2^2(8)$ and pairs of N2–H2···S1ⁱ/C2–H2B···S1ⁱ interactions, where rings of graph-set motif $R_2^1(7)$ are observed (Fig. 2, Table 1). As a

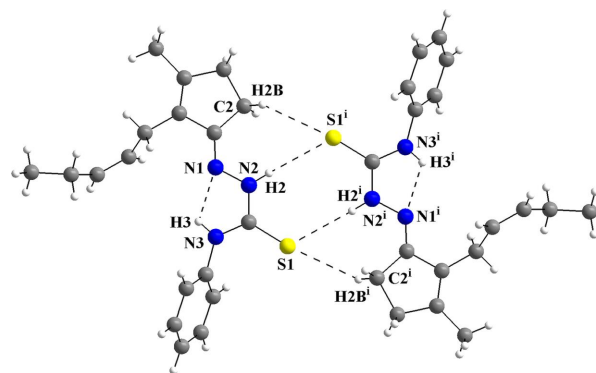


Figure 2
The molecular structure of the *cis*-jasmone 4-phenylthiosemicarbazone showing the intra- and intermolecular hydrogen-bond interactions as dashed lines. The molecules are linked into centrosymmetric dimers *via* pairs of N–H···S and C–H···S interactions, forming graph-set motifs $R_2^2(8)$ and $R_2^1(7)$. The N–H···N intramolecular interactions form rings with graph-set motif $S(5)$. [Symmetry code: (i) $-x + 1, -y, -z$.]

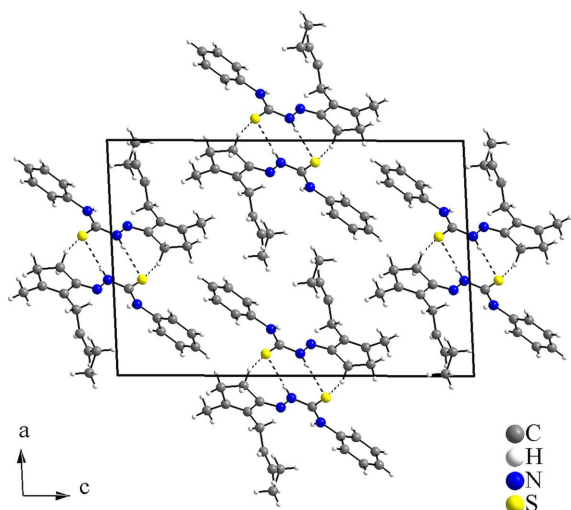


Figure 3
Crystal structure section of the title compound viewed along [010]. The hydrogen-bonding intermolecular interactions are drawn as dashed lines. The crystal structure resembles a zigzag motif when viewed from this direction.

feature of the dimeric structure, the sulfur atoms act as double acceptors and three rings with intermolecular hydrogen bonding are observed. No other strong intermolecular interactions can be suggested for the title compound due to the non-polar organic periphery and the steric effects of the phenyl ring and of the *cis*-jasmone fragment. Only weak interactions, *i.e.*, London dispersion forces, can be proposed. The crystal packing resembles a zigzag motif when viewed along [010] (Fig. 3).

For the title compound, the Hirshfeld surface analysis (Hirshfeld, 1977), the graphical representations and the two-dimensional Hirshfeld surface fingerprint (HSFP) were evaluated with the *Crystal Explorer* software (Wolff *et al.*, 2012). The graphical representation of the Hirshfeld surface (d_{norm}) is represented using a ball-and-stick model with transparency. In red, the locations of the strongest intermolecular contacts, *i.e.*, the regions around the H2 and S1 atoms (Fig. 4) are

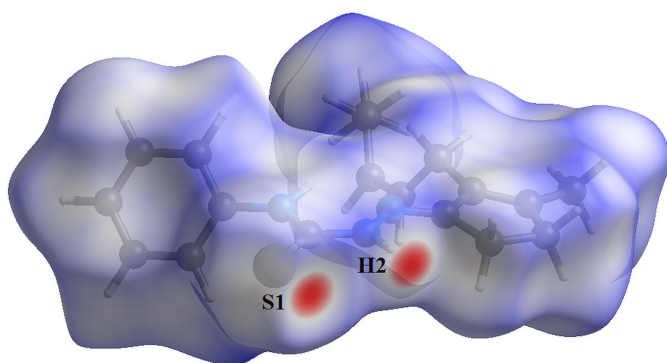


Figure 4
Hirshfeld surface graphical representation (d_{norm}) for the title compound. The molecule is drawn using a ball-and-stick model, the surface is drawn with transparency and the regions with strongest intermolecular interactions are shown in red and labeled. The figure is simplified for clarity. [d_{norm} range: -0.227 to 1.380].

indicated. These atoms are those involved in the $\text{H}\cdots\text{S}$ interactions showed in the previous figures (Figs. 2 and 3). The contributions to the crystal packing are shown as two-dimensional Hirshfeld surface fingerprint plots (HSFP) with cyan dots. The d_i (x -axis) and the d_e (y -axis) values are the closest internal and external distances from given points on the Hirshfeld surface contacts (in Å). The major contributions to the crystal packing amount to (a) $\text{H}\cdots\text{H} = 65.3\%$, (b) $\text{H}\cdots\text{C}/\text{C}\cdots\text{H} = 16.2\%$, (c) $\text{H}\cdots\text{S}/\text{S}\cdots\text{H} = 10.9\%$ and (d) $\text{H}\cdots\text{N}/\text{N}\cdots\text{H} = 5.5\%$ (Fig. 5).

To the best of our knowledge and from using database tools such as *SciFinder* (Chemical Abstracts Service, 2023) and the Cambridge Structural Database (CSD; Groom *et al.*, 2016), only the crystal structure of the non-substituted *cis*-jasmone thiosemicarbazone has been reported (Orsoni *et al.*, 2020). The terminal group of the thiosemicarbazones plays an essential role in the intermolecular interactions and the supramolecular arrangement, *e.g.*, the non-substituted form, which shows the NH_2 terminal group, leads to the building of mono-periodic hydrogen-bonded ribbons, while a phenyl ring attached to the terminal nitrogen atom leads to the formation of discrete dimeric units (Oliveira *et al.*, 2017). This molecular architecture is specially observed for compounds with a non-polar organic periphery and therefore, the tetralone 4-phenylthiosemicarbazone derivative (Oliveira *et al.*, 2014) was chosen for comparison with the title compound. As for the structure of the *cis*-jasmone 4-phenylthiosemicarbazone, an $\text{N3}-\text{H2N}\cdots\text{N2}$ intramolecular interaction is observed, with

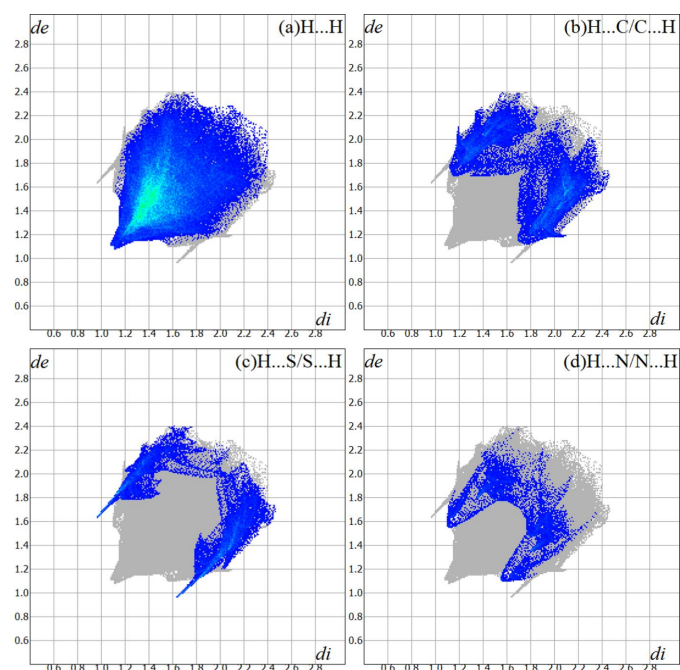


Figure 5
The Hirshfeld surface two-dimensional fingerprint plot (HSFP) for the title compound showing the intermolecular contacts in detail (cyan dots). The major contributions to the crystal cohesion amount to (a) $\text{H}\cdots\text{H} = 65.3\%$, (b) $\text{H}\cdots\text{C}/\text{C}\cdots\text{H} = 16.2\%$, (c) $\text{H}\cdots\text{S}/\text{S}\cdots\text{H} = 10.9\%$ and (d) $\text{H}\cdots\text{N}/\text{N}\cdots\text{H} = 5.5\%$. The d_i (x -axis) and the d_e (y -axis) values are the closest internal and external distances from given points on the Hirshfeld surface (in Å).

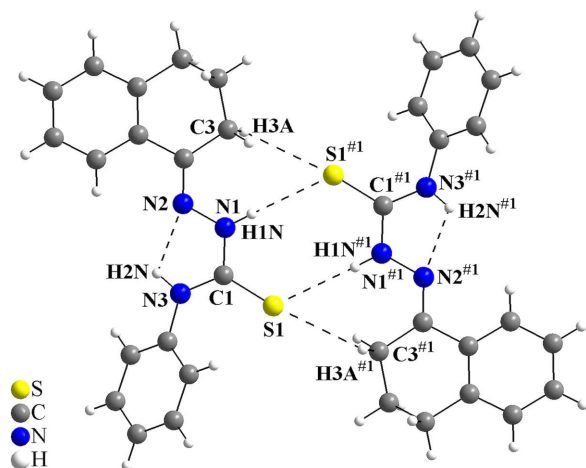


Figure 6
The molecular structure of the reference compound, tetralone 4-phenylthiosemicarbazone (Oliveira *et al.*, 2014), showing the intra- and intermolecular hydrogen-bond interactions drawn as dashed lines, which are quite similar to the title compound (Fig. 2). The molecules are linked into centrosymmetric dimers *via* pairs of N–H···S and C–H···S interactions, forming graph-set motifs of $R_2^2(8)$ and $R_2^1(7)$. The N–H···N intramolecular interactions, which form rings with graph-set motif $S(5)$, are also observed. [Symmetry code: (#1) $-x + 1, -y, -z + 1$.]

graph-set motif $S(5)$, and the thiosemicarbazone molecules are linked into centrosymmetric dimers *via* pairs of N1–H1N···S1^{#1} and C3–H3A···S1^{#1} interactions, forming hydrogen-bonded rings with graph-set motifs of $R_2^2(8)$ and $R_2^1(7)$. The sulfur atoms also act as double acceptors and, indeed, the intra and intermolecular hydrogen bonding in the structure of the tetralone 4-phenylthiosemicarbazone are quite similar to those of the title compound (for the dimeric structure and the symmetry code, see Fig. 6; for a structural comparison with the compound of this work, see: Fig. 2). In

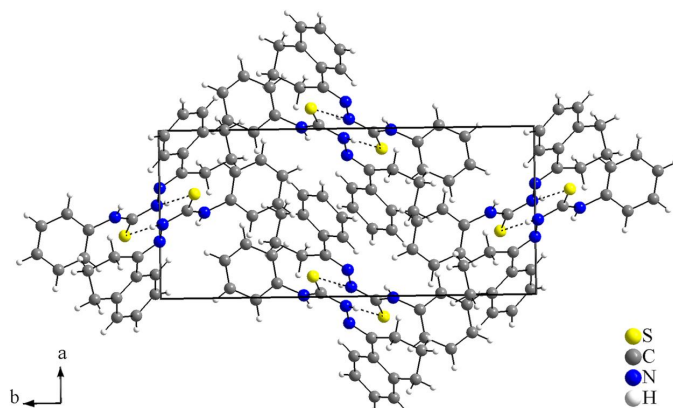


Figure 7
Crystal structure section of the comparison compound, tetralone 4-phenylthiosemicarbazone (Oliveira *et al.*, 2014), viewed along [001]. For this view, a zigzag motif of the discrete dimeric units can be suggested. It resembles the packing structure of the title compound (Fig. 3). Only the intermolecular N–H···S interactions are shown for clarity, drawn as dashed lines.

Table 2
Experimental details.

Crystal data	
Chemical formula	C ₁₈ H ₂₃ N ₃ S
<i>M_r</i>	313.45
Crystal system, space group	Monoclinic, $P2_1/n$
Temperature (K)	123
<i>a</i> , <i>b</i> , <i>c</i> (Å)	13.6565 (3), 5.8286 (2), 20.6721 (6)
β (°)	92.751 (2)
<i>V</i> (Å ³)	1643.57 (8)
<i>Z</i>	4
Radiation type	Mo $K\alpha$
μ (mm ⁻¹)	0.20
Crystal size (mm)	0.22 × 0.13 × 0.05
Data collection	
Diffractometer	Enraf–Nonius FR590 Kappa CCD
No. of measured, independent and observed [$I > 2\sigma(I)$] reflections	26959, 3751, 2857
R_{int}	0.064
($\sin \theta/\lambda$) _{max} (Å ⁻¹)	0.649
Refinement	
$R[F^2 > 2\sigma(F^2)]$, $wR(F^2)$, S	0.039, 0.095, 1.06
No. of reflections	3751
No. of parameters	202
H-atom treatment	H-atom parameters constrained
$\Delta\rho_{max}$, $\Delta\rho_{min}$ (e Å ⁻³)	0.28, -0.25

Computer programs: *COLLECT* (Nonius, 1998), *HKL DENZO* and *SCALEPACK* (Otwinowski & Minor, 1997), *SIR92* (Altomare *et al.*, 1999), *SHELXL2018/3* (Sheldrick, 2015), *DIAMOND* (Brandenburg, 2006), *CrystalExplorer* (Wolff *et al.*, 2012), *WinGX* (Farrugia, 2012), *pubCIF* (Westrip, 2010) and *enCIFer* (Allen *et al.*, 2004).

the crystal, viewed along [001], the tetralone 4-phenylthiosemicarbazone shows a also zigzag motif, resembling the packing structure of the title compound (Fig. 7).

Synthesis and crystallization

The starting materials are commercially available and were used without further purification. The synthesis was adapted from previously reported procedures (Freund & Schander, 1902; Oliveira *et al.*, 2014). The hydrochloric acid-catalyzed reaction between *cis*-jasmone (8 mmol) and 4-phenylthiosemicarbazide (8 mmol) in ethanol (80 ml) was refluxed for 6 h. After cooling and filtering, the title compound was obtained as precipitate, filtered off and washed with cold ethanol. Colorless single crystals suitable for X-ray diffraction were obtained in tetrahydrofuran by slow evaporation of the solvent.

Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2.

Acknowledgements

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full crystallographic data

IUCrData (2023). **8**, x230971 [https://doi.org/10.1107/S2414314623009719]

2-{3-Methyl-2-[(2*Z*)-pent-2-en-1-yl]cyclopent-2-en-1-ylidene}-*N*-phenylhydrazinecarbothioamide

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2-{3-Methyl-2-[(2*Z*)-pent-2-en-1-yl]cyclopent-2-en-1-ylidene}-*N*-phenylhydrazinecarbothioamide

Crystal data

C₁₈H₂₃N₃S

$M_r = 313.45$

Monoclinic, $P2_1/n$

$a = 13.6565$ (3) Å

$b = 5.8286$ (2) Å

$c = 20.6721$ (6) Å

$\beta = 92.751$ (2)°

$V = 1643.57$ (8) Å³

$Z = 4$

$F(000) = 672$

$D_x = 1.267$ Mg m⁻³

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

Cell parameters from 63134 reflections

$\theta = 2.9$ – 27.5 °

$\mu = 0.20$ mm⁻¹

$T = 123$ K

Fragment, colourless

$0.22 \times 0.13 \times 0.05$ mm

Data collection

Enraf–Nonius FR590 Kappa CCD diffractometer

Radiation source: sealed X-ray tube, Enraf–Nonius FR590

Detector resolution: 9 pixels mm⁻¹

CCD rotation images, thick slices, κ -goniostat scans

26959 measured reflections

3751 independent reflections

2857 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.064$

$\theta_{\text{max}} = 27.5$ °, $\theta_{\text{min}} = 3.0$ °

$h = -17 \rightarrow 17$

$k = -7 \rightarrow 7$

$l = -26 \rightarrow 26$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.039$

$wR(F^2) = 0.095$

$S = 1.06$

3751 reflections

202 parameters

0 restraints

Primary atom site location: structure-invariant direct methods

Hydrogen site location: inferred from neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0359P)^2 + 0.7368P]$

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

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

$\Delta\rho_{\text{max}} = 0.28$ e Å⁻³

$\Delta\rho_{\text{min}} = -0.25$ e Å⁻³

Extinction correction: SHELXL-2018/3 (Sheldrick 2015),

$F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0053 (9)

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.

Refinement. An absorption correction was not performed, as the crystal data analysis suggested that the absorption effects were not significant for the structure refinement. Hydrogen atoms were located in a difference-Fourier map, but were positioned with idealized geometry and refined isotropically using a riding model (HFIX command). Methyl H atoms were allowed to rotate but not to tip to best fit the experimental electron density. Thus, for the methyl H atoms [$U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{C})$], the C—H bond lengths were set to 0.96 Å. The other C—H bond lengths were also set according to the H atom neighbourhood [$U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{C})$]. For the phenyl H atoms and for the other H atoms attached to sp^2 -hybridized carbon atoms (C7 and C8), the C—H bond lengths were set 0.93 Å. For the H atoms of the —CH₂— fragments (C2, C3, C6 and C9), the C—H bond lengths were set to 0.97 Å. Finally, the N—H bond lengths [$U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}(\text{N})$] were set to 0.86 Å.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.38987 (11)	0.5560 (3)	−0.10461 (7)	0.0182 (3)
C2	0.45778 (11)	0.4298 (3)	−0.14766 (8)	0.0207 (3)
H2A	0.436900	0.272027	−0.154050	0.025*
H2B	0.524485	0.431089	−0.129277	0.025*
C3	0.45002 (12)	0.5646 (3)	−0.21191 (8)	0.0239 (4)
H3A	0.513393	0.625665	−0.222395	0.029*
H3B	0.425872	0.466784	−0.247139	0.029*
C4	0.37868 (11)	0.7559 (3)	−0.20041 (8)	0.0205 (3)
C5	0.34692 (11)	0.7526 (3)	−0.13988 (7)	0.0189 (3)
C6	0.27920 (11)	0.9155 (3)	−0.10790 (8)	0.0220 (3)
H6A	0.274508	1.055634	−0.133179	0.026*
H6B	0.307436	0.954690	−0.065378	0.026*
C7	0.17720 (12)	0.8219 (3)	−0.10048 (8)	0.0255 (4)
H7	0.165822	0.670153	−0.112577	0.031*
C8	0.10260 (12)	0.9383 (3)	−0.07815 (9)	0.0277 (4)
H8	0.043700	0.859385	−0.075799	0.033*
C9	0.10293 (12)	1.1834 (3)	−0.05628 (9)	0.0275 (4)
H9A	0.159387	1.260861	−0.072846	0.033*
H9B	0.109126	1.188208	−0.009357	0.033*
C10	0.01004 (13)	1.3102 (3)	−0.07920 (10)	0.0337 (4)
H10A	0.006058	1.315650	−0.125670	0.051*
H10B	0.011733	1.463645	−0.062278	0.051*
H10C	−0.046191	1.231343	−0.064102	0.051*
C11	0.35163 (13)	0.9202 (3)	−0.25364 (8)	0.0266 (4)
H11A	0.304018	1.027255	−0.239068	0.040*
H11B	0.324443	0.837326	−0.290397	0.040*
H11C	0.409031	1.001812	−0.265701	0.040*
C12	0.36853 (11)	0.2620 (3)	0.04217 (7)	0.0182 (3)
C13	0.24371 (11)	0.3876 (3)	0.11856 (7)	0.0189 (3)
C14	0.18756 (11)	0.1912 (3)	0.12727 (8)	0.0214 (3)
H14	0.190844	0.068671	0.098647	0.026*

C15	0.12685 (11)	0.1804 (3)	0.17895 (8)	0.0235 (4)
H15	0.088949	0.050178	0.184838	0.028*
C16	0.12210 (11)	0.3623 (3)	0.22203 (8)	0.0241 (4)
H16	0.081198	0.353730	0.256649	0.029*
C17	0.17823 (12)	0.5562 (3)	0.21342 (8)	0.0232 (4)
H17	0.175481	0.677598	0.242474	0.028*
C18	0.23886 (11)	0.5702 (3)	0.16133 (8)	0.0215 (3)
H18	0.276003	0.701443	0.155225	0.026*
N1	0.36550 (9)	0.5130 (2)	−0.04654 (6)	0.0189 (3)
N2	0.40538 (9)	0.3219 (2)	−0.01525 (6)	0.0195 (3)
H2	0.451694	0.244307	−0.031612	0.023*
N3	0.30211 (9)	0.4113 (2)	0.06367 (6)	0.0201 (3)
H3	0.294026	0.535658	0.041653	0.024*
S1	0.40757 (3)	0.02358 (7)	0.08139 (2)	0.02292 (12)

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0171 (8)	0.0187 (8)	0.0187 (8)	−0.0025 (6)	0.0011 (6)	−0.0009 (6)
C2	0.0207 (8)	0.0204 (8)	0.0210 (8)	0.0008 (6)	0.0031 (6)	0.0008 (6)
C3	0.0267 (9)	0.0242 (8)	0.0212 (8)	0.0006 (7)	0.0063 (7)	0.0007 (7)
C4	0.0193 (8)	0.0204 (8)	0.0216 (8)	−0.0030 (6)	0.0006 (6)	0.0015 (6)
C5	0.0180 (8)	0.0174 (8)	0.0212 (8)	−0.0024 (6)	0.0015 (6)	−0.0003 (6)
C6	0.0231 (8)	0.0177 (8)	0.0254 (9)	0.0008 (6)	0.0037 (7)	−0.0001 (7)
C7	0.0259 (9)	0.0192 (8)	0.0318 (10)	−0.0013 (7)	0.0047 (7)	−0.0009 (7)
C8	0.0236 (9)	0.0238 (9)	0.0361 (10)	−0.0022 (7)	0.0053 (7)	−0.0004 (7)
C9	0.0272 (9)	0.0252 (9)	0.0302 (9)	0.0012 (7)	0.0027 (7)	−0.0026 (7)
C10	0.0309 (10)	0.0251 (9)	0.0452 (12)	0.0030 (7)	0.0022 (8)	0.0007 (8)
C11	0.0294 (9)	0.0271 (9)	0.0231 (9)	−0.0017 (7)	−0.0006 (7)	0.0048 (7)
C12	0.0192 (8)	0.0180 (8)	0.0174 (8)	−0.0028 (6)	−0.0003 (6)	−0.0013 (6)
C13	0.0180 (8)	0.0207 (8)	0.0181 (8)	0.0030 (6)	0.0019 (6)	0.0030 (6)
C14	0.0234 (8)	0.0204 (8)	0.0205 (8)	−0.0004 (6)	0.0010 (7)	0.0003 (6)
C15	0.0213 (8)	0.0239 (8)	0.0255 (9)	−0.0034 (7)	0.0032 (7)	0.0051 (7)
C16	0.0203 (8)	0.0317 (9)	0.0207 (8)	0.0034 (7)	0.0042 (7)	0.0043 (7)
C17	0.0239 (8)	0.0244 (9)	0.0216 (8)	0.0048 (7)	0.0026 (7)	−0.0026 (6)
C18	0.0214 (8)	0.0199 (8)	0.0234 (8)	−0.0004 (6)	0.0027 (6)	0.0010 (6)
N1	0.0213 (7)	0.0165 (6)	0.0189 (7)	−0.0007 (5)	0.0010 (5)	0.0019 (5)
N2	0.0216 (7)	0.0183 (6)	0.0188 (7)	0.0029 (5)	0.0034 (5)	0.0011 (5)
N3	0.0240 (7)	0.0171 (6)	0.0195 (7)	0.0018 (5)	0.0057 (5)	0.0036 (5)
S1	0.0255 (2)	0.0203 (2)	0.0233 (2)	0.00340 (16)	0.00521 (16)	0.00501 (16)

Geometric parameters (Å, °)

C1—N1	1.286 (2)	C10—H10B	0.9600
C1—C5	1.466 (2)	C10—H10C	0.9600
C1—C2	1.507 (2)	C11—H11A	0.9600
C2—C3	1.542 (2)	C11—H11B	0.9600
C2—H2A	0.9700	C11—H11C	0.9600

C2—H2B	0.9700	C12—N3	1.348 (2)
C3—C4	1.507 (2)	C12—N2	1.357 (2)
C3—H3A	0.9700	C12—S1	1.6827 (16)
C3—H3B	0.9700	C13—C18	1.387 (2)
C4—C5	1.344 (2)	C13—C14	1.394 (2)
C4—C11	1.492 (2)	C13—N3	1.425 (2)
C5—C6	1.501 (2)	C14—C15	1.385 (2)
C6—C7	1.511 (2)	C14—H14	0.9300
C6—H6A	0.9700	C15—C16	1.388 (2)
C6—H6B	0.9700	C15—H15	0.9300
C7—C8	1.325 (2)	C16—C17	1.382 (2)
C7—H7	0.9300	C16—H16	0.9300
C8—C9	1.498 (2)	C17—C18	1.392 (2)
C8—H8	0.9300	C17—H17	0.9300
C9—C10	1.524 (2)	C18—H18	0.9300
C9—H9A	0.9700	N1—N2	1.3863 (18)
C9—H9B	0.9700	N2—H2	0.8600
C10—H10A	0.9600	N3—H3	0.8600
N1—C1—C5	120.09 (14)	C9—C10—H10B	109.5
N1—C1—C2	130.57 (14)	H10A—C10—H10B	109.5
C5—C1—C2	109.32 (13)	C9—C10—H10C	109.5
C1—C2—C3	103.95 (12)	H10A—C10—H10C	109.5
C1—C2—H2A	111.0	H10B—C10—H10C	109.5
C3—C2—H2A	111.0	C4—C11—H11A	109.5
C1—C2—H2B	111.0	C4—C11—H11B	109.5
C3—C2—H2B	111.0	H11A—C11—H11B	109.5
H2A—C2—H2B	109.0	C4—C11—H11C	109.5
C4—C3—C2	105.03 (13)	H11A—C11—H11C	109.5
C4—C3—H3A	110.7	H11B—C11—H11C	109.5
C2—C3—H3A	110.7	N3—C12—N2	113.96 (13)
C4—C3—H3B	110.7	N3—C12—S1	125.25 (12)
C2—C3—H3B	110.7	N2—C12—S1	120.78 (12)
H3A—C3—H3B	108.8	C18—C13—C14	120.27 (14)
C5—C4—C11	128.03 (15)	C18—C13—N3	118.60 (14)
C5—C4—C3	112.11 (14)	C14—C13—N3	121.01 (14)
C11—C4—C3	119.85 (14)	C15—C14—C13	119.37 (15)
C4—C5—C1	109.55 (14)	C15—C14—H14	120.3
C4—C5—C6	129.38 (15)	C13—C14—H14	120.3
C1—C5—C6	121.06 (14)	C14—C15—C16	120.55 (15)
C5—C6—C7	114.00 (13)	C14—C15—H15	119.7
C5—C6—H6A	108.8	C16—C15—H15	119.7
C7—C6—H6A	108.8	C17—C16—C15	119.91 (15)
C5—C6—H6B	108.8	C17—C16—H16	120.0
C7—C6—H6B	108.8	C15—C16—H16	120.0
H6A—C6—H6B	107.6	C16—C17—C18	120.12 (15)
C8—C7—C6	125.32 (15)	C16—C17—H17	119.9
C8—C7—H7	117.3	C18—C17—H17	119.9

C6—C7—H7	117.3	C13—C18—C17	119.78 (15)
C7—C8—C9	127.02 (16)	C13—C18—H18	120.1
C7—C8—H8	116.5	C17—C18—H18	120.1
C9—C8—H8	116.5	C1—N1—N2	118.59 (13)
C8—C9—C10	112.23 (15)	C12—N2—N1	117.45 (12)
C8—C9—H9A	109.2	C12—N2—H2	121.3
C10—C9—H9A	109.2	N1—N2—H2	121.3
C8—C9—H9B	109.2	C12—N3—C13	127.68 (13)
C10—C9—H9B	109.2	C12—N3—H3	116.2
H9A—C9—H9B	107.9	C13—N3—H3	116.2
C9—C10—H10A	109.5		
N1—C1—C2—C3	177.17 (16)	C18—C13—C14—C15	0.1 (2)
C5—C1—C2—C3	-1.15 (17)	N3—C13—C14—C15	-175.86 (14)
C1—C2—C3—C4	0.01 (16)	C13—C14—C15—C16	-0.4 (2)
C2—C3—C4—C5	1.26 (18)	C14—C15—C16—C17	0.1 (2)
C2—C3—C4—C11	-178.74 (14)	C15—C16—C17—C18	0.5 (2)
C11—C4—C5—C1	177.96 (15)	C14—C13—C18—C17	0.5 (2)
C3—C4—C5—C1	-2.03 (18)	N3—C13—C18—C17	176.55 (14)
C11—C4—C5—C6	-2.6 (3)	C16—C17—C18—C13	-0.8 (2)
C3—C4—C5—C6	177.45 (15)	C5—C1—N1—N2	178.18 (13)
N1—C1—C5—C4	-176.51 (14)	C2—C1—N1—N2	0.0 (2)
C2—C1—C5—C4	2.01 (18)	N3—C12—N2—N1	-5.15 (19)
N1—C1—C5—C6	4.0 (2)	S1—C12—N2—N1	176.26 (10)
C2—C1—C5—C6	-177.53 (13)	C1—N1—N2—C12	-171.07 (14)
C4—C5—C6—C7	104.92 (19)	N2—C12—N3—C13	173.62 (14)
C1—C5—C6—C7	-75.65 (19)	S1—C12—N3—C13	-7.9 (2)
C5—C6—C7—C8	-174.86 (17)	C18—C13—N3—C12	132.16 (16)
C6—C7—C8—C9	0.4 (3)	C14—C13—N3—C12	-51.8 (2)
C7—C8—C9—C10	138.6 (2)		

Hydrogen-bond geometry (\AA , $^\circ$)

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C2—H2B \cdots S1 ⁱ	0.97	2.96	3.4640 (16)	113
N2—H2 \cdots S1 ⁱ	0.86	2.72	3.5757 (13)	177
N3—H3 \cdots N1	0.86	2.11	2.5457 (18)	111

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