

Crystal structure of 2-[(2*E*)-2-methyl-3-phenylprop-2-en-1-ylidene]-*N*-phenylhydrazinecarboxamide

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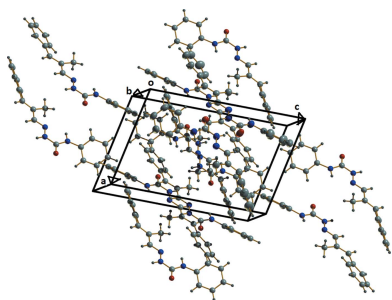
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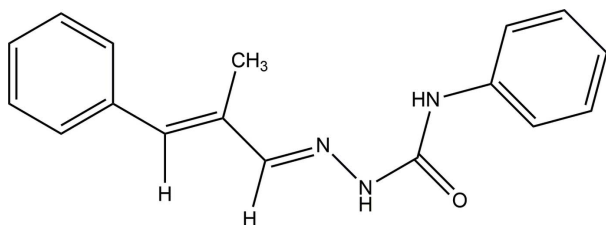
The title compound, C₁₇H₁₇N₃O, crystallizes with two independent molecules in the asymmetric unit. The semicarbazone moieties of these independent molecules (I and II) are essentially planar [maximum deviation of 0.042 (1) Å in molecule I and 0.041 (1) Å in molecule II], with the terminal phenyl rings twisted away from the mean plane of the semicarbazone moiety, making dihedral angles of 60.26 (8) and 28.76 (9)° in molecule I and 31.07 (9) and 35.45 (8)° in molecule II. The molecules both exhibit an *E* configuration with respect to the C=C and azomethine C=N bonds. In the crystal, two classical N—H···O hydrogen-bonding interactions are present between the two molecules, forming a centrosymmetric dimer, while a weak C—H···O non-classical hydrogen-bonding interaction, with a donor–acceptor distance of 3.476 (2) Å, interconnects two neighbouring centrosymmetric dimers to form a cage-like structure. These cage structures are interconnected by weak C—H···π interactions with an H···π distance of 2.790 Å, forming supramolecular chains along the *c*-axis direction.

1. Chemical context

Semicarbazones are oxygen and nitrogen contributor ligands whose significance lies in their versatility of molecular sequence, which allows diverse geometries to be obtained. Semicarbazones exhibit amido–iminol tautomerism in solution due to the interaction of solvent molecules, but generally exist in the amido form in the solid state. The FT–IR and NMR spectra of semicarbazones indicate the existence of a keto form in the solid state that can be confirmed by single crystal X-ray diffraction analysis (Kurup *et al.*, 2011; Sreekanth *et al.*, 2004). Biological properties linked to antimicrobial (Siji *et al.*, 2010) and antiparasitic (Soares *et al.*, 2011) effects make semicarbazones important ligands in coordination chemistry. Compared to Gentamycin, a commonly used antibiotic, *N*⁴-phenylsemicarbazone derivatives exhibit moderate antibacterial activity at higher concentrations and also show DNA cleavage properties (Layana *et al.*, 2016). Semicarbazones can function as brilliant ligands in a variety of metal ions (Kala *et al.*, 2007) and co-ordinate to metal ions either in neutral (Siji *et al.*, 2011) or in anionic forms (Reena *et al.*, 2008). Structural studies of many semicarbazones and *N*⁴-phenylsemicarbazones have been reported and some of them adopt an *E* configuration with respect to the azomethine double bond along with both inter- and intramolecular hydrogen-bonding interactions (Reena *et al.*, 2010; Layana *et al.*, 2014, 2018). Semicarbazones form complexes with a variety of structural



features such as monomer, dimer and one-dimensional polymers (Kunnath *et al.*, 2016). α -Methyl-*trans*-cinnamaldehyde, a precursor for the synthesis of α -methyl-*trans*-cinnamaldehyde- N^4 -phenylsemicarbazone, has significant antifungal activity and can self-couple and form complexes with some transition metals (Shreaz *et al.*, 2011). The diverse structural features and substantial biological applications have prompted us to synthesize a new semicarbazone derived from α -methyl-*trans*-cinnamaldehyde and N^4 -phenylsemicarbazide.



2. Structural commentary

The title compound crystallizes in the triclinic space group $P\bar{1}$ symmetry with two independent molecules, I and II, in the asymmetric unit (Fig. 1). The semicarbazone units in I and II are essentially planar, with maximum deviations from the least-squares plane of 0.042 (1) Å for N2 in molecule I and 0.041 (1) Å for N4 in molecule II. The terminal phenyl rings in both two molecules are twisted away from the semicarbazone mean plane, making dihedral angles of 60.26 (8) and 28.76 (9)° in molecule I and 31.07 (9) and 35.45 (8)° in molecule II. Both molecules exist in an *E* configuration with respect to the C=C and azomethine C=N bonds. The azomethine C=N and keto C=O bond lengths [1.273 (2) and 1.2269 (17) Å, respectively] in molecule I are shorter than those for molecule II [1.2766 (19) Å and 1.2302 (18) Å respectively]. In contrast, the C=N and C=O bond lengths reported for the two independent molecules of 2-benzoylpyridine semicarbazone are 1.294 (2) and 1.295 (2) Å and 1.2360 (19) and 1.2390 (19) Å respectively (de Lima *et al.*, 2008).

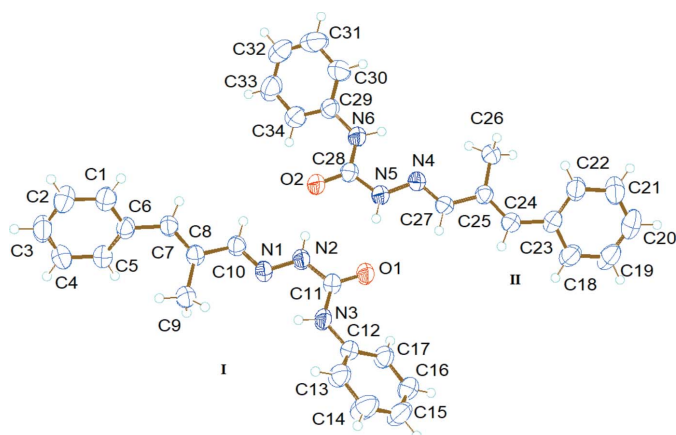


Figure 1
ORTEP diagram showing the two molecules in the asymmetric unit, with atom labels and 50% probability displacement ellipsoids.

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

Cg1 is the centroid of the C1–C6 ring.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
N5–H5'···O1	0.88 (1)	1.99 (1)	2.8639 (19)	174 (2)
N2–H2'···O2	0.88 (1)	1.93 (1)	2.808 (2)	176 (2)
N3–H3'···N1	0.87 (1)	2.13 (2)	2.6146 (18)	115 (1)
N6–H6'···N4	0.87 (1)	2.17 (2)	2.6261 (19)	112 (2)
C13–H13···O2 ⁱ	0.93	2.64	3.476 (2)	149
C32–H32···Cg1 ⁱⁱ	0.93	2.79	3.518 (2)	136

Symmetry codes: (i) $-x + 1, -y, -z + 1$; (ii) $-x + 1, -y, -z + 2$.

3. Supramolecular features

In the crystal, two classical and one non-classical hydrogen-bonding interactions are observed. Molecules I and II are linked into centrosymmetric dimers through N2–H2'···O2 and N5–H5'···O1 hydrogen bonds with $D\cdots A$ distances of 2.808 (2) Å, and 2.8639 (19) Å, respectively (Fig. 2, Table 1), while C13–H13···O2 interactions with a $D\cdots A$ distance of 3.476 (2) Å, interconnect adjacent dimers, creating cage-like structures that are linked by weak C–H··· π interactions into supramolecular chains along the *c*-axis direction (Fig. 3). No significant π – π interactions occur. The packing viewed along the *b* axis is shown in Fig. 4.

3.1. Database survey

The structure of the title compound has not previously been reported (CSD version 5.39, update of August 2018; Groom *et al.*, 2016). All geometric parameters in the title compound

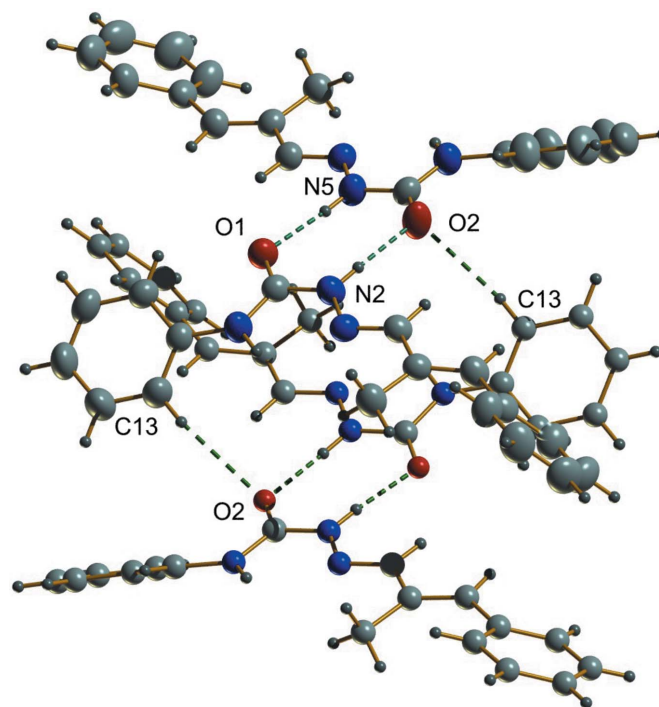


Figure 2
N–H···O hydrogen bonds and weak C–H···O intermolecular interactions (dashed lines) generating centrosymmetric dimers and a cage-like structure.

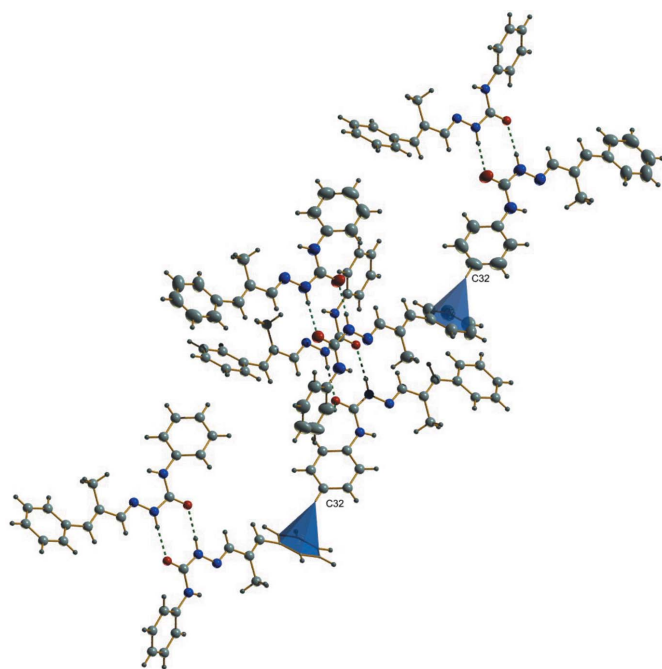


Figure 3
Weak C—H $\cdots\pi$ intermolecular interactions (solid cones), linking the dimeric cage-like structures into a chain along the *c* axis.

agree well with those reported in the literature with the C10—N1/C27—N4 [1.273 (2) and 1.2766 (19) Å], N1—N2/N4—N5 [1.3691 (17) and 1.3679 (18) Å] and C11—O1/C28—O2 [1.2269 (17) and 1.2302 (18) Å] bond distances being comparable to those in benzaldehyde-*N*⁴-phenylsemicarbazide

azone [1.273 (2), 1.369 (2) and 1.225 (2) Å; Layana *et al.*, 2014] and vanillin-*N*-phenylthiosemicarbazone [1.2726 (17), 1.3801 (15) and 1.2404 (15) Å; Layana *et al.*, 2016]

4. Synthesis and crystallization

Hot ethanolic solutions of *N*⁴-phenylsemicarbazide (0.1512 g, 1 mmol) and α -methyl-*trans*-cinnamaldehyde (0.14 ml, 1 mmol) were mixed and refluxed for about 4 h. Colourless block-shaped crystals of the title compound (yield 83%) were separated by filtration, washed with ethanol and dried over P₄O₁₀ *in vacuo*. Single crystals (m.p. 463±2 K) were obtained by slow evaporation of a 1:1 mixture of ethyl acetate and ethanol.

Analysis calculated: C, 73.03; H, 6.09, N, 15.04%. Found: C, 72.66; H, 6.32; N, 15.29%. Spectrometric data. FT-IR ν_{\max} (KBr, cm⁻¹): The spectrum of the title compound shows characteristic absorption bands of the main functional groups at IR (ν_{\max} , cm⁻¹): 3379 (⁴NH), 3192 (²NH), 1685 (C=O) 3072, 2960 (C—H aromatic), 1591 (C=N), 1029 (N—N). FT-Raman (cm⁻¹) 3055 (N—H), 1613 (C=O), 1577 (C=N), 1137 (N—N). ¹H NMR (400 MHz) (DMSO-*d*₆, ppm): δ_{H} 2.2 (s, 3H, methyl), 7–7.5 (*m*, 10H, Ar—H), 6.7 (s, 1H, methine), 7.7 (s, 1H, azomethine), 8.6 (s, 2H, amine), 10.6 (s, 1H, iminol H). ¹³C NMR (400 MHz) (DMSO-*d*₆, ppm): δ_{C} 135.2 (C6), 129.12 (C1 and C5), 128.4 (C2 and C4), 119.6 ppm (C3), 152.9 (C7), 146.4 (C8), 138.9 (C9), 136.5 (C10), 12.9 (C17), 134.3 (C11), 128.5 ppm (C12 and C16), 127.4 ppm (C13 and C15) and 122.4 ppm (C14). UV-visible (200–1000, nm): 268 (π – π^*), 342 (*n*– π^*).

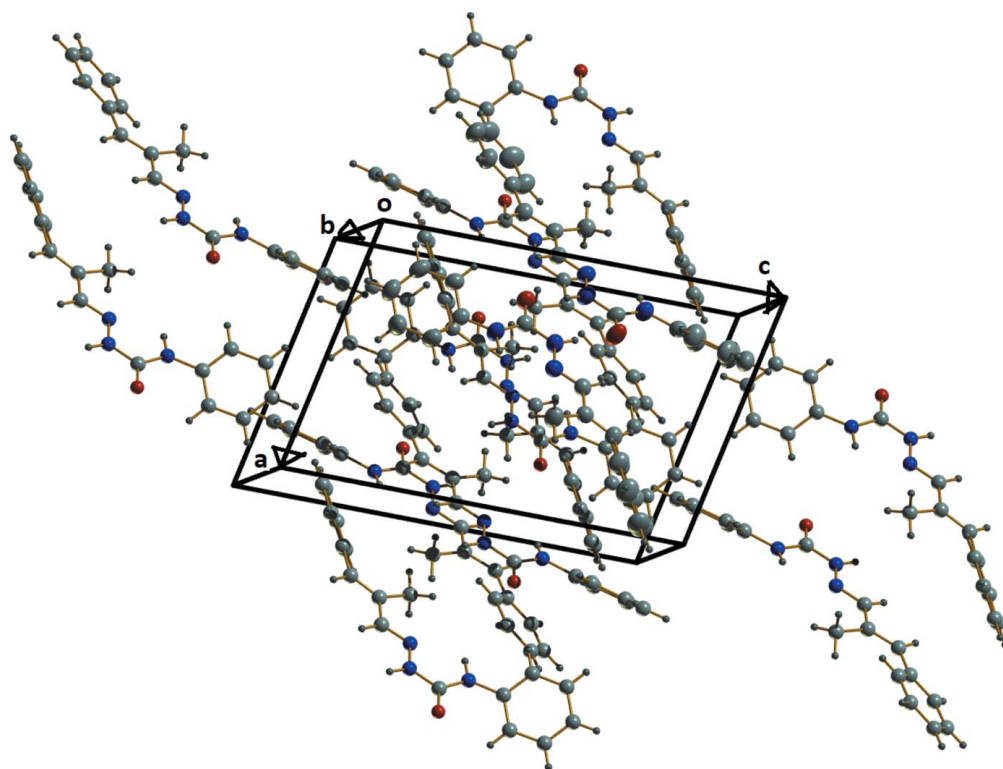


Figure 4
The packing viewed along the *b* axis.

Table 2
Experimental details.

Crystal data	
Chemical formula	C ₁₇ H ₁₇ N ₃ O
<i>M_r</i>	279.33
Crystal system, space group	Triclinic, <i>P</i> $\bar{1}$
Temperature (K)	296
<i>a</i> , <i>b</i> , <i>c</i> (Å)	10.2140 (6), 10.5133 (8), 15.3297 (10)
α , β , γ (°)	106.652 (3), 99.111 (3), 97.416 (4)
<i>V</i> (Å ³)	1530.51 (18)
<i>Z</i>	4
Radiation type	Mo <i>K</i> α
μ (mm ⁻¹)	0.08
Crystal size (mm)	0.60 × 0.50 × 0.50
Data collection	
Diffractometer	Bruker Kappa APEXII CCD
Absorption correction	Multi-scan (<i>SADABS</i> ; Bruker, 2004)
<i>T_{min}</i> , <i>T_{max}</i>	0.939, 0.948
No. of measured, independent and observed [<i>I</i> > 2σ(<i>I</i>)] reflections	12172, 7346, 4038
<i>R_{int}</i>	0.019
(sin θ /λ) _{max} (Å ⁻¹)	0.669
Refinement	
<i>R</i> [<i>F</i> ² > 2σ(<i>F</i> ²)], <i>wR</i> (<i>F</i> ²), <i>S</i>	0.051, 0.175, 0.95
No. of reflections	7346
No. of parameters	397
No. of restraints	4
H-atom treatment	H atoms treated by a mixture of independent and constrained refinement
$\Delta\rho_{\max}$, $\Delta\rho_{\min}$ (e Å ⁻³)	0.16, -0.23

Computer programs: *APEX2*, *SAINT* and *XPREP* (Bruker, 2004), *SHELXL2014* (Sheldrick, 2015), *SHELXS97* (Sheldrick, 2008), *ORTEP-3 for Windows* (Farrugia, 2012), *DIAMOND* (Brandenburg, 2010) and *pubCIF* (Westrip, 2010).

5. Refinement

Crystal data, data collection and structure refinement details are summarized in Table 2. Reflections ($\bar{1}\bar{1}1$) and (001) were omitted due to bad agreement. All hydrogen atoms bound to carbon atoms were positioned geometrically with C–H distances of 0.93–0.96 Å and refined as riding, with *U*_{iso}(H) = 1.2*U*_{eq}(C) or 1.5*U*_{eq}(C-methyl). The NH hydrogen atoms were located in a difference-Fourier map and refined with N–H restrained to 0.88±0.01 Å.

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spectroscopic studies and to the SAIF IIT Madras, Chennai, India for the FT–Raman spectroscopic data. They are grateful to Dr. M. R. Prathapachandra Kurup, Department of Applied Chemistry, Cochin University of Science & Technology, Kochi-22, India, for the use of the *DIAMOND* software.

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supporting information

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Computing details

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2* and *SAINT* (Bruker, 2004); data reduction: *SAINT* and *XPREP* (Bruker, 2004); program(s) used to solve structure: *SHELXL2014* (Sheldrick, 2015); program(s) used to refine structure: *SHELXS97* (Sheldrick, 2008); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 2012) and *DIAMOND* (Brandenburg, 2010); software used to prepare material for publication: *pubCIF* (Westrip, 2010).

2-[(2*E*)-2-Methyl-3-phenylprop-2-en-1-ylidene]-*N*-phenylhydrazinecarboxamide

Crystal data

$C_{17}H_{17}N_3O$	$Z = 4$
$M_r = 279.33$	$F(000) = 592$
Triclinic, $P\bar{1}$	$D_x = 1.212 \text{ Mg m}^{-3}$
$a = 10.2140 (6) \text{ \AA}$	Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
$b = 10.5133 (8) \text{ \AA}$	Cell parameters from 3309 reflections
$c = 15.3297 (10) \text{ \AA}$	$\theta = 2.6\text{--}27.7^\circ$
$\alpha = 106.652 (3)^\circ$	$\mu = 0.08 \text{ mm}^{-1}$
$\beta = 99.111 (3)^\circ$	$T = 296 \text{ K}$
$\gamma = 97.416 (4)^\circ$	Block, colorless
$V = 1530.51 (18) \text{ \AA}^3$	$0.60 \times 0.50 \times 0.50 \text{ mm}$

Data collection

Bruker Kappa APEXII CCD diffractometer	12172 measured reflections
Radiation source: fine-focus sealed tube	7346 independent reflections
Graphite monochromator	4038 reflections with $I > 2\sigma(I)$
ω and ϕ scan	$R_{\text{int}} = 0.019$
Absorption correction: multi-scan (SADABS; Bruker, 2004)	$\theta_{\text{max}} = 28.4^\circ$, $\theta_{\text{min}} = 2.2^\circ$
$T_{\text{min}} = 0.939$, $T_{\text{max}} = 0.948$	$h = -13 \rightarrow 12$
	$k = -13 \rightarrow 13$
	$l = -20 \rightarrow 15$

Refinement

Refinement on F^2	Hydrogen site location: mixed
Least-squares matrix: full	H atoms treated by a mixture of independent and constrained refinement
$R[F^2 > 2\sigma(F^2)] = 0.051$	$w = 1/[\sigma^2(F_o^2) + (0.1017P)^2]$
$wR(F^2) = 0.175$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 0.95$	$(\Delta/\sigma)_{\text{max}} < 0.001$
7346 reflections	$\Delta\rho_{\text{max}} = 0.16 \text{ e \AA}^{-3}$
397 parameters	$\Delta\rho_{\text{min}} = -0.23 \text{ e \AA}^{-3}$
4 restraints	

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.88173 (19)	-0.1561 (2)	0.82683 (13)	0.0630 (5)
H1	0.8824	-0.0637	0.8438	0.076*
C2	0.9866 (2)	-0.2034 (3)	0.86846 (15)	0.0723 (6)
H2	1.0576	-0.1430	0.9125	0.087*
C3	0.9863 (2)	-0.3387 (3)	0.84506 (16)	0.0753 (6)
H3	1.0568	-0.3706	0.8732	0.090*
C4	0.8822 (2)	-0.4271 (2)	0.78030 (15)	0.0734 (6)
H4	0.8818	-0.5194	0.7647	0.088*
C5	0.7773 (2)	-0.3802 (2)	0.73766 (13)	0.0613 (5)
H5	0.7073	-0.4415	0.6933	0.074*
C6	0.77527 (16)	-0.2438 (2)	0.76005 (12)	0.0497 (4)
C7	0.66419 (17)	-0.18955 (19)	0.71894 (12)	0.0502 (4)
H7	0.6304	-0.1251	0.7604	0.060*
C8	0.60596 (15)	-0.22156 (18)	0.62933 (11)	0.0442 (4)
C9	0.64815 (19)	-0.3180 (2)	0.55182 (13)	0.0611 (5)
H9A	0.7350	-0.3360	0.5739	0.092*
H9B	0.6534	-0.2795	0.5025	0.092*
H9C	0.5834	-0.4008	0.5290	0.092*
C10	0.49813 (15)	-0.15284 (18)	0.60452 (11)	0.0448 (4)
H10	0.4637	-0.0979	0.6512	0.054*
C11	0.29530 (15)	-0.10122 (17)	0.41634 (11)	0.0438 (4)
C12	0.32548 (15)	-0.18810 (17)	0.25412 (11)	0.0447 (4)
C13	0.42849 (18)	-0.2055 (2)	0.20670 (13)	0.0649 (6)
H13	0.5143	-0.2049	0.2382	0.078*
C14	0.4051 (2)	-0.2239 (3)	0.11315 (14)	0.0850 (8)
H14	0.4754	-0.2355	0.0815	0.102*
C15	0.2788 (2)	-0.2254 (3)	0.06550 (14)	0.0777 (7)
H15	0.2634	-0.2373	0.0020	0.093*
C16	0.17664 (19)	-0.2093 (2)	0.11236 (13)	0.0640 (5)
H16	0.0907	-0.2110	0.0804	0.077*
C17	0.19852 (16)	-0.1905 (2)	0.20641 (12)	0.0541 (5)
H17	0.1278	-0.1794	0.2377	0.065*
C18	-0.33204 (19)	0.2586 (2)	0.20723 (13)	0.0665 (6)
H18	-0.3167	0.1721	0.1801	0.080*
C19	-0.4276 (2)	0.3093 (3)	0.15904 (15)	0.0787 (7)
H19	-0.4753	0.2573	0.0999	0.094*
C20	-0.4518 (2)	0.4367 (3)	0.19863 (16)	0.0793 (7)
H20	-0.5171	0.4706	0.1670	0.095*
C21	-0.3796 (2)	0.5128 (3)	0.28446 (17)	0.0769 (6)

H21	-0.3953	0.5993	0.3110	0.092*
C22	-0.2836 (2)	0.4633 (2)	0.33250 (14)	0.0643 (5)
H22	-0.2347	0.5174	0.3908	0.077*
C23	-0.25859 (16)	0.3344 (2)	0.29541 (12)	0.0507 (4)
C24	-0.15906 (16)	0.27357 (19)	0.34168 (12)	0.0498 (4)
H24	-0.1223	0.2101	0.3017	0.060*
C25	-0.11244 (15)	0.29476 (17)	0.43239 (11)	0.0431 (4)
C26	-0.15538 (17)	0.38927 (19)	0.51113 (12)	0.0516 (4)
H26A	-0.0976	0.4761	0.5297	0.077*
H26B	-0.1493	0.3543	0.5627	0.077*
H26C	-0.2469	0.3982	0.4916	0.077*
C27	-0.01521 (15)	0.21447 (17)	0.45568 (11)	0.0451 (4)
H27	0.0122	0.1537	0.4079	0.054*
C28	0.17376 (15)	0.13993 (19)	0.63940 (12)	0.0474 (4)
C29	0.17182 (15)	0.24127 (19)	0.80442 (12)	0.0482 (4)
C30	0.1829 (2)	0.3675 (2)	0.86669 (14)	0.0740 (6)
H30	0.1683	0.4397	0.8452	0.089*
C31	0.2159 (2)	0.3874 (3)	0.96110 (15)	0.0878 (7)
H31	0.2219	0.4729	1.0029	0.105*
C32	0.23933 (19)	0.2836 (3)	0.99329 (15)	0.0733 (6)
H32	0.2616	0.2974	1.0569	0.088*
C33	0.2302 (2)	0.1594 (3)	0.93208 (15)	0.0766 (7)
H33	0.2476	0.0882	0.9540	0.092*
C34	0.1954 (2)	0.1371 (2)	0.83737 (14)	0.0675 (5)
H34	0.1880	0.0510	0.7961	0.081*
N1	0.44999 (12)	-0.16681 (14)	0.51975 (9)	0.0447 (3)
N2	0.34882 (14)	-0.09748 (16)	0.50394 (10)	0.0485 (4)
N3	0.35427 (14)	-0.17255 (17)	0.34979 (10)	0.0536 (4)
N4	0.03382 (12)	0.22481 (14)	0.53988 (9)	0.0455 (3)
N5	0.12237 (14)	0.14139 (16)	0.55249 (10)	0.0516 (4)
N6	0.13105 (14)	0.22397 (17)	0.70885 (10)	0.0538 (4)
O1	0.20160 (11)	-0.04300 (13)	0.40141 (8)	0.0525 (3)
O2	0.25451 (13)	0.06620 (15)	0.65140 (8)	0.0648 (4)
H3'	0.4260 (12)	-0.1964 (17)	0.3742 (11)	0.049 (5)*
H5'	0.1507 (17)	0.0898 (17)	0.5059 (9)	0.060 (6)*
H6'	0.080 (2)	0.276 (2)	0.6905 (15)	0.087 (7)*
H2'	0.3176 (16)	-0.0501 (17)	0.5509 (9)	0.055 (5)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0670 (11)	0.0636 (14)	0.0591 (11)	0.0160 (10)	0.0032 (9)	0.0238 (10)
C2	0.0595 (11)	0.0850 (18)	0.0701 (13)	0.0139 (11)	-0.0047 (10)	0.0299 (13)
C3	0.0660 (12)	0.0986 (19)	0.0719 (14)	0.0403 (12)	0.0071 (11)	0.0358 (14)
C4	0.0886 (14)	0.0696 (15)	0.0705 (14)	0.0405 (12)	0.0118 (12)	0.0262 (12)
C5	0.0664 (11)	0.0615 (14)	0.0568 (11)	0.0221 (9)	0.0036 (9)	0.0204 (10)
C6	0.0519 (9)	0.0611 (12)	0.0446 (10)	0.0185 (8)	0.0118 (8)	0.0253 (9)
C7	0.0561 (10)	0.0558 (12)	0.0463 (10)	0.0227 (8)	0.0123 (8)	0.0209 (9)

C8	0.0415 (8)	0.0493 (10)	0.0467 (10)	0.0124 (7)	0.0108 (7)	0.0198 (8)
C9	0.0615 (11)	0.0734 (14)	0.0521 (11)	0.0290 (10)	0.0109 (8)	0.0186 (10)
C10	0.0455 (8)	0.0494 (11)	0.0446 (9)	0.0136 (7)	0.0116 (7)	0.0192 (8)
C11	0.0416 (8)	0.0477 (10)	0.0457 (9)	0.0108 (7)	0.0081 (7)	0.0197 (8)
C12	0.0453 (9)	0.0457 (10)	0.0419 (9)	0.0117 (7)	0.0044 (7)	0.0128 (8)
C13	0.0478 (10)	0.0906 (16)	0.0518 (11)	0.0219 (10)	0.0054 (8)	0.0141 (11)
C14	0.0620 (12)	0.138 (2)	0.0514 (12)	0.0239 (13)	0.0164 (10)	0.0196 (14)
C15	0.0725 (13)	0.113 (2)	0.0420 (11)	0.0223 (12)	0.0025 (10)	0.0188 (12)
C16	0.0545 (10)	0.0749 (15)	0.0532 (11)	0.0158 (9)	-0.0066 (9)	0.0134 (10)
C17	0.0432 (9)	0.0663 (13)	0.0519 (11)	0.0134 (8)	0.0044 (8)	0.0185 (9)
C18	0.0621 (11)	0.0881 (17)	0.0470 (11)	0.0220 (10)	0.0057 (9)	0.0169 (11)
C19	0.0634 (12)	0.120 (2)	0.0508 (12)	0.0199 (13)	-0.0005 (9)	0.0295 (13)
C20	0.0599 (12)	0.116 (2)	0.0783 (16)	0.0254 (13)	0.0060 (11)	0.0555 (16)
C21	0.0739 (13)	0.0754 (16)	0.0897 (16)	0.0222 (11)	0.0032 (12)	0.0422 (14)
C22	0.0677 (12)	0.0597 (14)	0.0642 (12)	0.0129 (9)	-0.0033 (9)	0.0257 (11)
C23	0.0463 (9)	0.0648 (13)	0.0449 (10)	0.0096 (8)	0.0081 (7)	0.0240 (9)
C24	0.0504 (9)	0.0522 (11)	0.0458 (10)	0.0135 (8)	0.0078 (7)	0.0133 (8)
C25	0.0403 (8)	0.0432 (10)	0.0439 (9)	0.0064 (7)	0.0046 (7)	0.0137 (8)
C26	0.0530 (9)	0.0535 (12)	0.0508 (10)	0.0179 (8)	0.0109 (8)	0.0163 (9)
C27	0.0445 (8)	0.0455 (10)	0.0431 (9)	0.0114 (7)	0.0069 (7)	0.0103 (8)
C28	0.0397 (8)	0.0527 (11)	0.0461 (10)	0.0114 (7)	0.0022 (7)	0.0120 (8)
C29	0.0404 (8)	0.0587 (12)	0.0454 (9)	0.0135 (7)	0.0081 (7)	0.0147 (9)
C30	0.0992 (15)	0.0679 (15)	0.0585 (13)	0.0323 (12)	0.0211 (11)	0.0154 (11)
C31	0.1139 (19)	0.0819 (18)	0.0553 (14)	0.0197 (14)	0.0196 (13)	0.0005 (13)
C32	0.0642 (12)	0.104 (2)	0.0472 (12)	0.0100 (12)	0.0097 (9)	0.0199 (13)
C33	0.0860 (15)	0.0902 (19)	0.0625 (14)	0.0214 (13)	0.0101 (11)	0.0380 (14)
C34	0.0845 (13)	0.0593 (14)	0.0558 (12)	0.0146 (10)	0.0066 (10)	0.0174 (10)
N1	0.0417 (7)	0.0494 (9)	0.0485 (8)	0.0147 (6)	0.0088 (6)	0.0214 (7)
N2	0.0490 (8)	0.0593 (10)	0.0444 (8)	0.0231 (7)	0.0120 (6)	0.0207 (7)
N3	0.0499 (8)	0.0735 (11)	0.0438 (8)	0.0295 (7)	0.0071 (6)	0.0216 (8)
N4	0.0417 (7)	0.0472 (9)	0.0465 (8)	0.0127 (6)	0.0045 (6)	0.0136 (7)
N5	0.0512 (8)	0.0588 (10)	0.0441 (9)	0.0247 (7)	0.0042 (7)	0.0119 (8)
N6	0.0547 (8)	0.0642 (11)	0.0449 (8)	0.0263 (7)	0.0072 (7)	0.0159 (8)
O1	0.0492 (6)	0.0641 (9)	0.0501 (7)	0.0244 (6)	0.0089 (5)	0.0218 (6)
O2	0.0679 (8)	0.0774 (10)	0.0485 (7)	0.0406 (7)	0.0013 (6)	0.0124 (7)

Geometric parameters (Å, °)

C1—C2	1.381 (2)	C19—C20	1.375 (3)
C1—C6	1.389 (3)	C19—H19	0.9300
C1—H1	0.9300	C20—C21	1.361 (3)
C2—C3	1.362 (3)	C20—H20	0.9300
C2—H2	0.9300	C21—C22	1.378 (2)
C3—C4	1.364 (3)	C21—H21	0.9300
C3—H3	0.9300	C22—C23	1.386 (3)
C4—C5	1.385 (2)	C22—H22	0.9300
C4—H4	0.9300	C23—C24	1.464 (2)
C5—C6	1.379 (3)	C24—C25	1.340 (2)

C5—H5	0.9300	C24—H24	0.9300
C6—C7	1.469 (2)	C25—C27	1.450 (2)
C7—C8	1.334 (2)	C25—C26	1.493 (2)
C7—H7	0.9300	C26—H26A	0.9600
C8—C10	1.453 (2)	C26—H26B	0.9600
C8—C9	1.487 (2)	C26—H26C	0.9600
C9—H9A	0.9600	C27—N4	1.2766 (19)
C9—H9B	0.9600	C27—H27	0.9300
C9—H9C	0.9600	C28—O2	1.2302 (18)
C10—N1	1.273 (2)	C28—N6	1.346 (2)
C10—H10	0.9300	C28—N5	1.358 (2)
C11—O1	1.2269 (17)	C29—C34	1.363 (3)
C11—N2	1.355 (2)	C29—C30	1.373 (3)
C11—N3	1.356 (2)	C29—N6	1.410 (2)
C12—C13	1.373 (2)	C30—C31	1.380 (3)
C12—C17	1.378 (2)	C30—H30	0.9300
C12—N3	1.406 (2)	C31—C32	1.354 (3)
C13—C14	1.368 (3)	C31—H31	0.9300
C13—H13	0.9300	C32—C33	1.353 (3)
C14—C15	1.374 (3)	C32—H32	0.9300
C14—H14	0.9300	C33—C34	1.382 (3)
C15—C16	1.359 (3)	C33—H33	0.9300
C15—H15	0.9300	C34—H34	0.9300
C16—C17	1.375 (3)	N1—N2	1.3691 (17)
C16—H16	0.9300	N2—H2'	0.881 (9)
C17—H17	0.9300	N3—H3'	0.868 (9)
C18—C19	1.383 (3)	N4—N5	1.3679 (18)
C18—C23	1.390 (3)	N5—H5'	0.879 (9)
C18—H18	0.9300	N6—H6'	0.872 (9)
C2—C1—C6	121.3 (2)	C21—C20—H20	120.2
C2—C1—H1	119.4	C19—C20—H20	120.2
C6—C1—H1	119.4	C20—C21—C22	120.8 (2)
C3—C2—C1	120.0 (2)	C20—C21—H21	119.6
C3—C2—H2	120.0	C22—C21—H21	119.6
C1—C2—H2	120.0	C21—C22—C23	121.1 (2)
C2—C3—C4	119.84 (18)	C21—C22—H22	119.5
C2—C3—H3	120.1	C23—C22—H22	119.5
C4—C3—H3	120.1	C22—C23—C18	117.31 (16)
C3—C4—C5	120.4 (2)	C22—C23—C24	124.73 (16)
C3—C4—H4	119.8	C18—C23—C24	117.95 (18)
C5—C4—H4	119.8	C25—C24—C23	130.34 (16)
C6—C5—C4	120.90 (19)	C25—C24—H24	114.8
C6—C5—H5	119.6	C23—C24—H24	114.8
C4—C5—H5	119.6	C24—C25—C27	116.69 (15)
C5—C6—C1	117.52 (16)	C24—C25—C26	125.97 (15)
C5—C6—C7	122.88 (17)	C27—C25—C26	117.30 (14)
C1—C6—C7	119.57 (17)	C25—C26—H26A	109.5

C8—C7—C6	127.81 (16)	C25—C26—H26B	109.5
C8—C7—H7	116.1	H26A—C26—H26B	109.5
C6—C7—H7	116.1	C25—C26—H26C	109.5
C7—C8—C10	118.17 (15)	H26A—C26—H26C	109.5
C7—C8—C9	124.61 (15)	H26B—C26—H26C	109.5
C10—C8—C9	117.15 (14)	N4—C27—C25	121.85 (15)
C8—C9—H9A	109.5	N4—C27—H27	119.1
C8—C9—H9B	109.5	C25—C27—H27	119.1
H9A—C9—H9B	109.5	O2—C28—N6	123.87 (15)
C8—C9—H9C	109.5	O2—C28—N5	120.66 (16)
H9A—C9—H9C	109.5	N6—C28—N5	115.48 (14)
H9B—C9—H9C	109.5	C34—C29—C30	118.95 (17)
N1—C10—C8	120.96 (15)	C34—C29—N6	122.71 (17)
N1—C10—H10	119.5	C30—C29—N6	118.29 (17)
C8—C10—H10	119.5	C29—C30—C31	120.2 (2)
O1—C11—N2	120.94 (15)	C29—C30—H30	119.9
O1—C11—N3	124.61 (14)	C31—C30—H30	119.9
N2—C11—N3	114.45 (13)	C32—C31—C30	120.5 (2)
C13—C12—C17	119.33 (16)	C32—C31—H31	119.8
C13—C12—N3	117.83 (13)	C30—C31—H31	119.8
C17—C12—N3	122.81 (15)	C33—C32—C31	119.5 (2)
C14—C13—C12	120.14 (16)	C33—C32—H32	120.3
C14—C13—H13	119.9	C31—C32—H32	120.3
C12—C13—H13	119.9	C32—C33—C34	120.8 (2)
C13—C14—C15	120.64 (19)	C32—C33—H33	119.6
C13—C14—H14	119.7	C34—C33—H33	119.6
C15—C14—H14	119.7	C29—C34—C33	120.1 (2)
C16—C15—C14	119.16 (18)	C29—C34—H34	120.0
C16—C15—H15	120.4	C33—C34—H34	120.0
C14—C15—H15	120.4	C10—N1—N2	116.30 (14)
C15—C16—C17	120.91 (16)	C11—N2—N1	120.63 (14)
C15—C16—H16	119.5	C11—N2—H2'	119.3 (11)
C17—C16—H16	119.5	N1—N2—H2'	120.1 (11)
C16—C17—C12	119.81 (17)	C11—N3—C12	127.30 (13)
C16—C17—H17	120.1	C11—N3—H3'	111.3 (11)
C12—C17—H17	120.1	C12—N3—H3'	120.4 (11)
C19—C18—C23	121.3 (2)	C27—N4—N5	116.14 (14)
C19—C18—H18	119.4	C28—N5—N4	120.26 (15)
C23—C18—H18	119.4	C28—N5—H5'	117.5 (12)
C20—C19—C18	119.9 (2)	N4—N5—H5'	122.2 (12)
C20—C19—H19	120.0	C28—N6—C29	125.66 (14)
C18—C19—H19	120.0	C28—N6—H6'	113.7 (15)
C21—C20—C19	119.53 (19)	C29—N6—H6'	120.3 (15)
C6—C1—C2—C3	-0.8 (3)	C22—C23—C24—C25	29.1 (3)
C1—C2—C3—C4	0.1 (3)	C18—C23—C24—C25	-152.1 (2)
C2—C3—C4—C5	0.5 (4)	C23—C24—C25—C27	178.40 (17)
C3—C4—C5—C6	-0.4 (3)	C23—C24—C25—C26	0.9 (3)

C4—C5—C6—C1	-0.2 (3)	C24—C25—C27—N4	-178.73 (16)
C4—C5—C6—C7	-177.99 (19)	C26—C25—C27—N4	-1.0 (2)
C2—C1—C6—C5	0.8 (3)	C34—C29—C30—C31	-0.8 (3)
C2—C1—C6—C7	178.68 (18)	N6—C29—C30—C31	176.69 (19)
C5—C6—C7—C8	-47.8 (3)	C29—C30—C31—C32	0.9 (4)
C1—C6—C7—C8	134.4 (2)	C30—C31—C32—C33	-0.1 (4)
C6—C7—C8—C10	-179.13 (17)	C31—C32—C33—C34	-1.0 (3)
C6—C7—C8—C9	-2.4 (3)	C30—C29—C34—C33	-0.2 (3)
C7—C8—C10—N1	171.39 (16)	N6—C29—C34—C33	-177.57 (17)
C9—C8—C10—N1	-5.6 (3)	C32—C33—C34—C29	1.1 (3)
C17—C12—C13—C14	0.6 (3)	C8—C10—N1—N2	-179.88 (14)
N3—C12—C13—C14	178.7 (2)	O1—C11—N2—N1	177.56 (15)
C12—C13—C14—C15	-0.1 (4)	N3—C11—N2—N1	-2.8 (2)
C13—C14—C15—C16	-0.4 (4)	C10—N1—N2—C11	177.09 (15)
C14—C15—C16—C17	0.6 (4)	O1—C11—N3—C12	3.3 (3)
C15—C16—C17—C12	-0.1 (3)	N2—C11—N3—C12	-176.34 (17)
C13—C12—C17—C16	-0.4 (3)	C13—C12—N3—C11	149.8 (2)
N3—C12—C17—C16	-178.43 (18)	C17—C12—N3—C11	-32.2 (3)
C23—C18—C19—C20	-0.4 (3)	C25—C27—N4—N5	178.64 (15)
C18—C19—C20—C21	1.2 (4)	O2—C28—N5—N4	-179.26 (16)
C19—C20—C21—C22	-0.7 (4)	N6—C28—N5—N4	0.2 (2)
C20—C21—C22—C23	-0.7 (3)	C27—N4—N5—C28	-176.31 (16)
C21—C22—C23—C18	1.4 (3)	O2—C28—N6—C29	0.5 (3)
C21—C22—C23—C24	-179.73 (19)	N5—C28—N6—C29	-179.02 (16)
C19—C18—C23—C22	-0.9 (3)	C34—C29—N6—C28	-37.1 (3)
C19—C18—C23—C24	-179.82 (19)	C30—C29—N6—C28	145.5 (2)

Hydrogen-bond geometry (Å, °)

Cg1 is the centroid of the C1—C6 ring.

<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>
N5—H5'...O1	0.88 (1)	1.99 (1)	2.8639 (19)	174 (2)
N2—H2'...O2	0.88 (1)	1.93 (1)	2.808 (2)	176 (2)
N3—H3'...N1	0.87 (1)	2.13 (2)	2.6146 (18)	115 (1)
N6—H6'...N4	0.87 (1)	2.17 (2)	2.6261 (19)	112 (2)
C13—H13...O2 ⁱ	0.93	2.64	3.476 (2)	149
C32—H32...Cg1 ⁱⁱ	0.93	2.79	3.518 (2)	136

Symmetry codes: (i) $-x+1, -y, -z+1$; (ii) $-x+1, -y, -z+2$.