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A new polymorph of *N*-(2- $\{N'$ -[(1*E*)-2-hydroxybenzylidene]hydrazinylcarbonyl}-phenyl)benzamide

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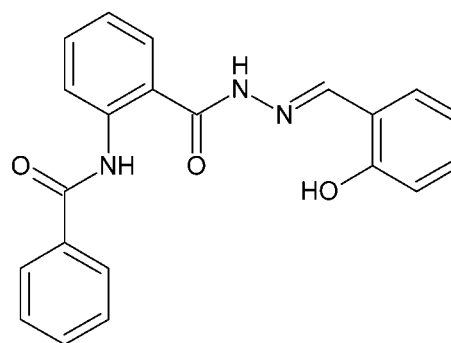
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Key indicators: single-crystal X-ray study; $T = 150$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.046; wR factor = 0.107; data-to-parameter ratio = 20.0.

The title compound, $\text{C}_{21}\text{H}_{17}\text{N}_3\text{O}_3$, is a new polymorph of an already published structure [Shashidhar *et al.* (2006). *Acta Cryst. E* **62**, o4473–o4475]. The previously reported structure crystallizes in the monoclinic space group $C2/c$, whereas the structure reported here is in the tetragonal space group $I4_1/a$. The bond lengths and angles are similar in both structures. The molecule adopts an extended conformation *via* intramolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{N}$ hydrogen bonds; the terminal phenyl ring and the hydroxylphenyl ring are twisted with respect to the central benzene ring by 44.43 (7) and 21.99 (8)°, respectively. In the crystal, molecules are linked by $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds, weak $\text{C}-\text{H}\cdots\text{O}$ hydrogen bonds and weak $\text{C}-\text{H}\cdots\pi$ interactions into a three-dimensional supra-molecular network.

Related literature

For different medicinal functions of hydrazide–hydrazone compounds, see: Bharti *et al.* (2010); Loncle *et al.* (2004); Garoufalas *et al.* (2002); Vicini *et al.* (2002); Sondhi *et al.* (2006); Kaymakçioğlu & Rollas (2002); Rahman *et al.* (2005); Ragavendran *et al.* (2007); Çakır *et al.* (2001); Terzioglu & Gursoy (2003); Vicini *et al.* (2009). For a monoclinic polymorph of the title compound, see: Shashidhar *et al.* (2006).



Experimental

Crystal data

$\text{C}_{21}\text{H}_{17}\text{N}_3\text{O}_3$
 $M_r = 359.38$
 Tetragonal, $I4_1/a$
 $a = 26.7145$ (14) Å
 $c = 10.1160$ (5) Å
 $V = 7219.4$ (8) Å³
 $Z = 16$
 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 150$ K
 $0.22 \times 0.19 \times 0.15$ mm

Data collection

Bruker SMART APEX CCD diffractometer
 Absorption correction: multi-scan (SADABS; Bruker, 2013)
 $T_{\min} = 0.80$, $T_{\max} = 0.99$
 37068 measured reflections
 4868 independent reflections
 3625 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.049$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.107$
 $S = 1.04$
 4868 reflections
 244 parameters
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.23$ e Å⁻³
 $\Delta\rho_{\min} = -0.21$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

Cg1 is the centroid of the C1–C6 phenyl ring.

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
$\text{N}2-\text{H}2\text{A}\cdots\text{O}1^{\text{i}}$	0.91	2.02	2.9181 (15)	171
$\text{N}3-\text{H}3\text{A}\cdots\text{O}2$	0.91	1.85	2.6373 (16)	143
$\text{O}3-\text{H}3\text{B}\cdots\text{N}1$	0.84	1.85	2.6052 (16)	148
$\text{C}4-\text{H}4\cdots\text{O}2^{\text{ii}}$	0.95	2.60	3.3388 (19)	135
$\text{C}12-\text{H}12\cdots\text{O}1^{\text{i}}$	0.95	2.47	3.2319 (17)	137
$\text{C}18-\text{H}18\cdots\text{O}3^{\text{iii}}$	0.95	2.53	3.417 (2)	155
$\text{C}17-\text{H}17\cdots\text{Cg}1^{\text{i}}$	0.95	2.73	3.6561 (18)	166

Symmetry codes: (i) $-y + \frac{5}{4}, x - \frac{1}{4}, z - \frac{1}{4}$; (ii) $-x + 1, -y + \frac{1}{2}, z$; (iii) $y + \frac{3}{4}, -x + \frac{3}{4}, z - \frac{1}{4}$.

Data collection: APEX2 (Bruker, 2013); cell refinement: SAINT (Bruker, 2013); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Sheldrick, 2008); program(s) used to refine structure: SHELXL2014 (Sheldrick, 2008); molecular graphics: DIAMOND (Brandenburg & Putz, 2012); software used to prepare material for publication: SHELXTL.

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Supporting information for this paper is available from the IUCr electronic archives (Reference: XU5789).

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supplementary materials

Acta Cryst. (2014). E70, o645–o646 [doi:10.1107/S1600536814010010]

A new polymorph of *N*-(2-*N'*-[(1*E*)-2-hydroxybenzylidene]hydrazinecarbonyl)-phenyl)benzamide

Shaaban K. Mohamed, Joel T. Mague, Mehmet Akkurt, Herman Potgieter and Mustafa R. Albayati

1. Comment

In many reports, hydrazide-hydrazone compounds are considered to be good candidates for different pharmaceutical applications such as anti-bacterial (Bharti *et al.*, 2010), antifugal (Loncle *et al.*, 2004), antimicrobial (Garoufalas *et al.*, 2002; Vicini *et al.*, 2002), anti-inflammatory (Sondhi *et al.*, 2006), anti-malarial and anti-tuberculosic activities (Kaymakçioğlu & Rollas, 2002; Rahman *et al.*, 2005) as well as anticonvulsant agents (Ragavendran *et al.*, 2007; Çakır *et al.*, 2001). In addition, hydrazide-hydrazones were reported to elicit anti-cancer (Terzioglu & Gursoy, 2003) and anti-HIV properties (Vicini *et al.*, 2009) and hence they have gained an important place in medicinal chemistry. As part of our study to obtain novel hydrazide-hydrazones with a wide spectrum of pharmaceutical properties we report herein the synthesis of the title compound as an example of a series of hydrazide-hydrazones.

The title compound, (I), is a tetragonal polymorph of the previously reported crystal structure which crystallizes in the monoclinic space group *C2/c* (Shashidhar *et al.*, 2006). The relative arrangement of the molecules in (I) is different from that previously reported.

The "extended" conformation of the title molecule (I) is largely determined by the intramolecular N—H \cdots O and O—H \cdots N hydrogen bonds (Table 1 and Fig. 1). The dihedral angle between the central 6-membered ring (C8–C13) and the phenyl ring (C1–C6) of the benzamide group is 44.43 (7) $^\circ$ while that with the phenol ring (C16–C21) is 21.99 (8) $^\circ$. The molecular packing of (I) is stabilized by intermolecular N—H \cdots O hydrogen bonds a weak C—H \cdots π interaction (Table 1 and Fig. 2).

2. Experimental

In 25 ml of ethanol, a mixture of 237 mg (1 mmol) 3-amino-2-phenylquinazolin-4(3*H*)-one, 122 mg (1 mmol) of salicaldehyde and a catalytic amount of glacial acetic acid was stirred and refluxed for 6hr. The mixture was cooled and the separated solid was recrystallized from ethanol to afford the pure product as pale-yellow crystals suitable for X-ray diffraction. m.p. 497 – 501 K.

3. Refinement

H-atoms attached to carbon were placed in calculated positions (C—H = 0.95 Å) while those attached to nitrogen and oxygen were placed in locations derived from a difference map and their coordinates adjusted to give N—H = 0.91 and O—H = 0.84 Å. All were included as riding contributions with isotropic displacement parameters 1.2 times those of the attached atoms.

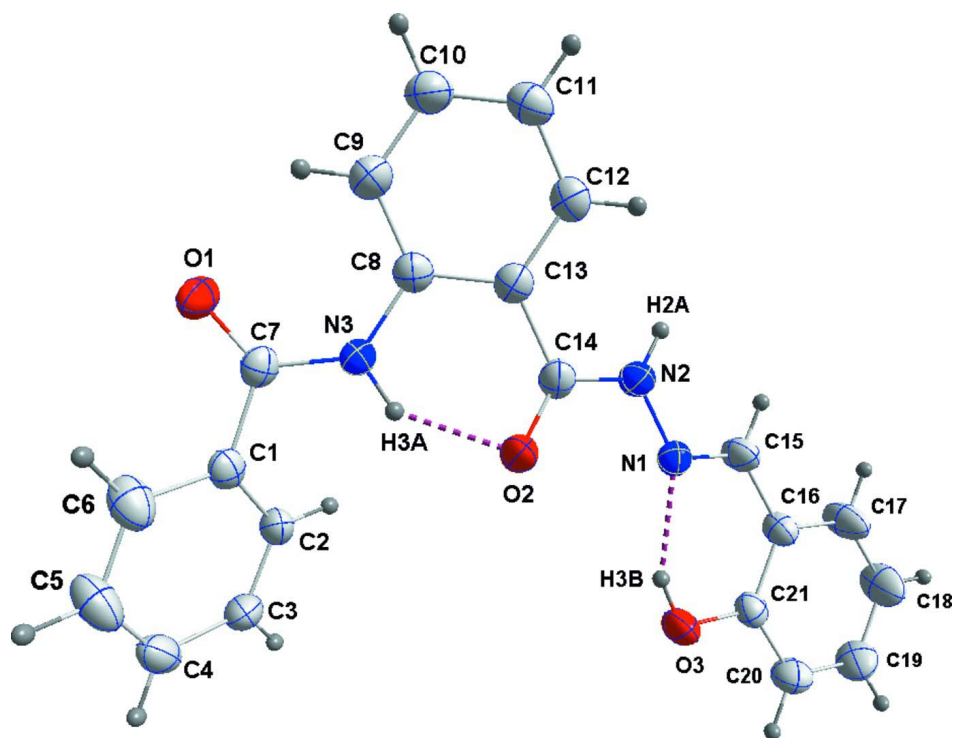


Figure 1

Perspective view of the title molecule with 50% probability ellipsoids. Intramolecular hydrogen bonds are shown by dotted lines.

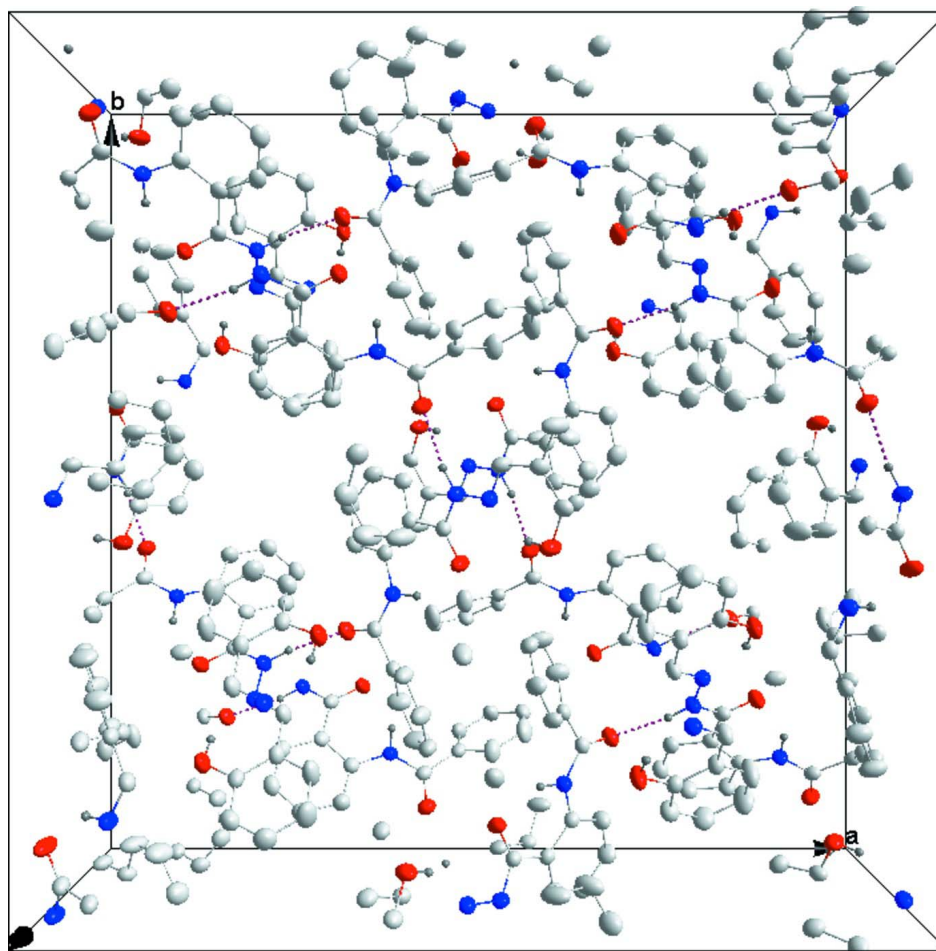


Figure 2

Packing viewed down the *c* axis with intermolecular N—H···O hydrogen bonds shown with dotted lines.

***N*-(2-{*N'*-[(1*E*)-2-Hydroxybenzylidene]hydrazinecarbonyl}phenyl)benzamide**

Crystal data

$C_{21}H_{17}N_3O_3$

$M_r = 359.38$

Tetragonal, $I4_1/a$

Hall symbol: $-I\ 4ad$

$a = 26.7145$ (14) Å

$c = 10.1160$ (5) Å

$V = 7219.4$ (8) Å³

$Z = 16$

$F(000) = 3008$

$D_x = 1.323$ Mg m⁻³

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

Cell parameters from 9997 reflections

$\theta = 2.2$ – 29.4°

$\mu = 0.09$ mm⁻¹

$T = 150$ K

Block, pale-yellow

$0.22 \times 0.19 \times 0.15$ mm

Data collection

Bruker SMART APEX CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 8.3660 pixels mm⁻¹

φ and ω scans

Absorption correction: multi-scan

(*SADABS*; Bruker, 2013)

$T_{\min} = 0.80$, $T_{\max} = 0.99$

37068 measured reflections

4868 independent reflections

3625 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.049$
 $\theta_{\text{max}} = 29.5^\circ$, $\theta_{\text{min}} = 2.2^\circ$
 $h = -36 \rightarrow 36$

$k = -36 \rightarrow 36$
 $l = -13 \rightarrow 13$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.046$
 $wR(F^2) = 0.107$
 $S = 1.04$
 4868 reflections
 244 parameters
 0 restraints

Hydrogen site location: inferred from neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0318P)^2 + 5.367P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\text{max}} = 0.001$
 $\Delta\rho_{\text{max}} = 0.23 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.21 \text{ e } \text{\AA}^{-3}$

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement on F^2 for ALL reflections except those flagged by the user for potential systematic errors. Weighted R -factors wR and all goodnesses of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The observed criterion of $F^2 > \sigma(F^2)$ is used only for calculating $-R$ -factor-obs *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

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

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.56564 (4)	0.41265 (4)	0.31994 (10)	0.0379 (3)
O2	0.65352 (4)	0.27173 (4)	0.11479 (12)	0.0444 (4)
O3	0.67750 (4)	0.17105 (4)	-0.15220 (12)	0.0461 (4)
N1	0.72994 (4)	0.23746 (4)	-0.02418 (12)	0.0323 (3)
N2	0.73567 (4)	0.27577 (4)	0.06534 (12)	0.0343 (4)
N3	0.61447 (4)	0.35295 (4)	0.22122 (12)	0.0327 (3)
C1	0.52568 (5)	0.35160 (5)	0.18534 (14)	0.0307 (4)
C2	0.52601 (5)	0.33600 (5)	0.05509 (16)	0.0372 (4)
C3	0.48315 (6)	0.31576 (6)	-0.00106 (17)	0.0416 (5)
C4	0.44051 (6)	0.30914 (5)	0.07402 (17)	0.0407 (5)
C5	0.44026 (6)	0.32438 (7)	0.20375 (18)	0.0521 (6)
C6	0.48221 (6)	0.34632 (7)	0.25975 (16)	0.0457 (5)
C7	0.57029 (5)	0.37577 (5)	0.24815 (14)	0.0302 (4)
C8	0.66281 (5)	0.36385 (5)	0.26991 (14)	0.0310 (4)
C9	0.67193 (6)	0.40316 (6)	0.35685 (16)	0.0401 (5)
C10	0.71960 (6)	0.41150 (6)	0.40559 (16)	0.0419 (5)
C11	0.75900 (6)	0.38104 (6)	0.36873 (15)	0.0389 (5)
C12	0.75057 (5)	0.34228 (5)	0.28113 (14)	0.0342 (4)
C13	0.70295 (5)	0.33311 (5)	0.22863 (14)	0.0298 (4)
C14	0.69475 (5)	0.29129 (5)	0.13280 (15)	0.0325 (4)
C15	0.76871 (5)	0.22376 (6)	-0.08880 (15)	0.0378 (5)
C16	0.76600 (5)	0.18314 (5)	-0.18336 (14)	0.0333 (4)
C17	0.80939 (6)	0.16779 (7)	-0.24892 (18)	0.0525 (6)
C18	0.80889 (7)	0.12900 (8)	-0.33838 (18)	0.0565 (6)
C19	0.76431 (6)	0.10478 (7)	-0.36461 (16)	0.0476 (5)

C20	0.72072 (6)	0.11928 (6)	-0.30239 (16)	0.0416 (5)
C21	0.72114 (5)	0.15829 (5)	-0.21138 (14)	0.0320 (4)
H2	0.55560	0.33910	0.00390	0.0450*
H2A	0.76580	0.29150	0.06880	0.0410*
H3	0.48320	0.30640	-0.09170	0.0500*
H3A	0.61360	0.32350	0.17590	0.0390*
H3B	0.68340	0.19490	-0.10030	0.0550*
H4	0.41160	0.29420	0.03630	0.0490*
H5	0.41100	0.31980	0.25570	0.0620*
H6	0.48130	0.35770	0.34870	0.0550*
H9	0.64520	0.42450	0.38300	0.0480*
H10	0.72530	0.43840	0.46500	0.0500*
H11	0.79160	0.38670	0.40330	0.0470*
H12	0.77770	0.32140	0.25590	0.0410*
H15	0.79970	0.24040	-0.07460	0.0450*
H17	0.84000	0.18460	-0.23130	0.0630*
H18	0.83880	0.11900	-0.38160	0.0680*
H19	0.76370	0.07790	-0.42610	0.0570*
H20	0.69020	0.10250	-0.32180	0.0500*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0422 (6)	0.0323 (5)	0.0391 (6)	0.0078 (4)	-0.0023 (4)	-0.0068 (4)
O2	0.0287 (5)	0.0375 (6)	0.0669 (8)	-0.0050 (4)	0.0089 (5)	-0.0155 (5)
O3	0.0265 (5)	0.0485 (6)	0.0633 (8)	-0.0034 (4)	0.0024 (5)	-0.0142 (5)
N1	0.0298 (6)	0.0300 (6)	0.0370 (6)	0.0003 (4)	0.0016 (5)	-0.0009 (5)
N2	0.0283 (6)	0.0334 (6)	0.0412 (7)	-0.0039 (5)	0.0043 (5)	-0.0057 (5)
N3	0.0301 (6)	0.0276 (5)	0.0404 (7)	0.0001 (4)	0.0010 (5)	-0.0059 (5)
C1	0.0298 (6)	0.0250 (6)	0.0372 (7)	0.0026 (5)	0.0028 (6)	0.0027 (5)
C2	0.0300 (7)	0.0372 (7)	0.0444 (9)	-0.0025 (6)	0.0084 (6)	-0.0096 (6)
C3	0.0362 (8)	0.0401 (8)	0.0485 (9)	-0.0030 (6)	0.0035 (7)	-0.0153 (7)
C4	0.0318 (7)	0.0340 (7)	0.0564 (10)	-0.0049 (6)	0.0011 (7)	0.0010 (7)
C5	0.0329 (8)	0.0766 (12)	0.0467 (10)	-0.0075 (8)	0.0087 (7)	0.0117 (9)
C6	0.0364 (8)	0.0676 (11)	0.0332 (8)	-0.0010 (7)	0.0047 (6)	0.0041 (7)
C7	0.0334 (7)	0.0261 (6)	0.0310 (7)	0.0022 (5)	0.0029 (5)	0.0023 (5)
C8	0.0320 (7)	0.0292 (6)	0.0318 (7)	-0.0013 (5)	0.0013 (5)	0.0026 (5)
C9	0.0398 (8)	0.0375 (8)	0.0430 (9)	0.0015 (6)	-0.0007 (7)	-0.0071 (7)
C10	0.0448 (8)	0.0431 (8)	0.0379 (8)	-0.0041 (7)	-0.0041 (7)	-0.0067 (7)
C11	0.0355 (7)	0.0481 (9)	0.0330 (8)	-0.0050 (6)	-0.0040 (6)	0.0011 (6)
C12	0.0322 (7)	0.0374 (7)	0.0331 (7)	-0.0010 (5)	0.0033 (6)	0.0039 (6)
C13	0.0317 (7)	0.0278 (6)	0.0298 (7)	-0.0026 (5)	0.0042 (5)	0.0035 (5)
C14	0.0303 (7)	0.0275 (6)	0.0398 (8)	-0.0007 (5)	0.0047 (6)	0.0022 (6)
C15	0.0265 (7)	0.0440 (8)	0.0428 (9)	-0.0055 (6)	0.0024 (6)	-0.0054 (7)
C16	0.0270 (6)	0.0403 (8)	0.0325 (7)	-0.0001 (5)	-0.0004 (5)	-0.0015 (6)
C17	0.0299 (8)	0.0763 (12)	0.0514 (10)	-0.0046 (8)	0.0050 (7)	-0.0231 (9)
C18	0.0407 (9)	0.0768 (13)	0.0520 (11)	0.0056 (8)	0.0063 (8)	-0.0239 (9)
C19	0.0531 (10)	0.0507 (9)	0.0390 (9)	0.0032 (7)	-0.0011 (7)	-0.0118 (7)
C20	0.0407 (8)	0.0441 (8)	0.0399 (9)	-0.0047 (7)	-0.0063 (7)	-0.0033 (7)
C21	0.0284 (6)	0.0339 (7)	0.0336 (7)	0.0020 (5)	-0.0023 (5)	0.0049 (6)

Geometric parameters (Å, °)

O1—C7	1.2303 (17)	C12—C13	1.4002 (19)
O2—C14	1.2326 (17)	C13—C14	1.495 (2)
O3—C21	1.3542 (17)	C15—C16	1.448 (2)
O3—H3B	0.8400	C16—C21	1.3990 (19)
N1—N2	1.3751 (16)	C16—C17	1.397 (2)
N1—C15	1.2783 (18)	C17—C18	1.376 (3)
N2—C14	1.3537 (18)	C18—C19	1.381 (3)
N3—C7	1.3560 (17)	C19—C20	1.379 (2)
N3—C8	1.4125 (17)	C20—C21	1.391 (2)
N2—H2A	0.9100	C2—H2	0.9500
N3—H3A	0.9100	C3—H3	0.9500
C1—C7	1.4970 (19)	C4—H4	0.9500
C1—C2	1.382 (2)	C5—H5	0.9500
C1—C6	1.391 (2)	C6—H6	0.9500
C2—C3	1.388 (2)	C9—H9	0.9500
C3—C4	1.381 (2)	C10—H10	0.9500
C4—C5	1.374 (2)	C11—H11	0.9500
C5—C6	1.386 (2)	C12—H12	0.9500
C8—C13	1.4137 (19)	C15—H15	0.9500
C8—C9	1.391 (2)	C17—H17	0.9500
C9—C10	1.384 (2)	C18—H18	0.9500
C10—C11	1.382 (2)	C19—H19	0.9500
C11—C12	1.381 (2)	C20—H20	0.9500
C21—O3—H3B	108.00	C16—C17—C18	121.69 (15)
N2—N1—C15	117.37 (11)	C17—C18—C19	119.18 (17)
N1—N2—C14	118.04 (11)	C18—C19—C20	120.59 (17)
C7—N3—C8	129.27 (12)	C19—C20—C21	120.38 (15)
C14—N2—H2A	124.00	O3—C21—C20	118.32 (13)
N1—N2—H2A	118.00	C16—C21—C20	119.78 (13)
C7—N3—H3A	118.00	O3—C21—C16	121.90 (12)
C8—N3—H3A	112.00	C1—C2—H2	120.00
C6—C1—C7	118.60 (13)	C3—C2—H2	120.00
C2—C1—C6	119.38 (13)	C2—C3—H3	120.00
C2—C1—C7	121.98 (12)	C4—C3—H3	120.00
C1—C2—C3	120.16 (14)	C3—C4—H4	120.00
C2—C3—C4	120.37 (15)	C5—C4—H4	120.00
C3—C4—C5	119.44 (15)	C4—C5—H5	120.00
C4—C5—C6	120.79 (15)	C6—C5—H5	120.00
C1—C6—C5	119.79 (15)	C1—C6—H6	120.00
N3—C7—C1	114.44 (12)	C5—C6—H6	120.00
O1—C7—N3	124.51 (13)	C8—C9—H9	120.00
O1—C7—C1	121.04 (12)	C10—C9—H9	120.00
N3—C8—C13	118.08 (12)	C9—C10—H10	120.00
N3—C8—C9	122.42 (12)	C11—C10—H10	120.00
C9—C8—C13	119.50 (13)	C10—C11—H11	120.00
C8—C9—C10	120.53 (14)	C12—C11—H11	120.00
C9—C10—C11	120.67 (15)	C11—C12—H12	119.00

C10—C11—C12	119.37 (14)	C13—C12—H12	119.00
C11—C12—C13	121.51 (13)	N1—C15—H15	120.00
C8—C13—C14	120.96 (12)	C16—C15—H15	120.00
C8—C13—C12	118.39 (12)	C16—C17—H17	119.00
C12—C13—C14	120.65 (12)	C18—C17—H17	119.00
O2—C14—N2	121.15 (13)	C17—C18—H18	120.00
O2—C14—C13	122.92 (13)	C19—C18—H18	120.00
N2—C14—C13	115.94 (11)	C18—C19—H19	120.00
N1—C15—C16	120.78 (13)	C20—C19—H19	120.00
C15—C16—C21	122.15 (12)	C19—C20—H20	120.00
C17—C16—C21	118.38 (13)	C21—C20—H20	120.00
C15—C16—C17	119.47 (13)		
C15—N1—N2—C14	178.84 (13)	C9—C8—C13—C12	2.4 (2)
N2—N1—C15—C16	178.99 (12)	C9—C8—C13—C14	-178.53 (13)
N1—N2—C14—O2	0.0 (2)	C8—C9—C10—C11	0.1 (2)
N1—N2—C14—C13	-179.85 (11)	C9—C10—C11—C12	0.7 (2)
C8—N3—C7—O1	2.7 (2)	C10—C11—C12—C13	0.0 (2)
C8—N3—C7—C1	-175.82 (13)	C11—C12—C13—C8	-1.6 (2)
C7—N3—C8—C9	-1.3 (2)	C11—C12—C13—C14	179.33 (13)
C7—N3—C8—C13	178.23 (13)	C8—C13—C14—O2	-22.0 (2)
C6—C1—C2—C3	0.7 (2)	C8—C13—C14—N2	157.86 (13)
C7—C1—C2—C3	-177.16 (13)	C12—C13—C14—O2	157.06 (14)
C2—C1—C6—C5	1.7 (2)	C12—C13—C14—N2	-23.07 (19)
C7—C1—C6—C5	179.57 (15)	N1—C15—C16—C17	-177.66 (15)
C2—C1—C7—O1	137.04 (15)	N1—C15—C16—C21	2.1 (2)
C2—C1—C7—N3	-44.39 (18)	C15—C16—C17—C18	179.19 (16)
C6—C1—C7—O1	-40.8 (2)	C21—C16—C17—C18	-0.5 (2)
C6—C1—C7—N3	137.75 (14)	C15—C16—C21—O3	0.5 (2)
C1—C2—C3—C4	-2.7 (2)	C15—C16—C21—C20	-179.54 (14)
C2—C3—C4—C5	2.3 (2)	C17—C16—C21—O3	-179.83 (14)
C3—C4—C5—C6	0.1 (3)	C17—C16—C21—C20	0.2 (2)
C4—C5—C6—C1	-2.1 (3)	C16—C17—C18—C19	0.4 (3)
N3—C8—C9—C10	177.83 (14)	C17—C18—C19—C20	0.2 (3)
C13—C8—C9—C10	-1.7 (2)	C18—C19—C20—C21	-0.5 (3)
N3—C8—C13—C12	-177.16 (12)	C19—C20—C21—O3	-179.64 (14)
N3—C8—C13—C14	1.9 (2)	C19—C20—C21—C16	0.4 (2)

Hydrogen-bond geometry (Å, °)

Cg1 is the centroid of the C1—C6 phenyl ring.

<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>
N2—H2 <i>A</i> \cdots O1 ⁱ	0.91	2.02	2.9181 (15)	171
N3—H3 <i>A</i> \cdots O2	0.91	1.85	2.6373 (16)	143
O3—H3 <i>B</i> \cdots N1	0.84	1.85	2.6052 (16)	148
C4—H4 \cdots O2 ⁱⁱ	0.95	2.60	3.3388 (19)	135
C9—H9 \cdots O1	0.95	2.24	2.8751 (19)	123
C12—H12 \cdots O1 ⁱ	0.95	2.47	3.2319 (17)	137

C18—H18 \cdots O3 ⁱⁱⁱ	0.95	2.53	3.417 (2)	155
C17—H17 \cdots Cg1 ⁱ	0.95	2.73	3.6561 (18)	166

Symmetry codes: (i) $-y+5/4, x-1/4, z-1/4$; (ii) $-x+1, -y+1/2, z$; (iii) $y+3/4, -x+3/4, z-1/4$.