

(E)-1-Phenylbutan-2-one (2,4-dinitrophenyl)hydrazone

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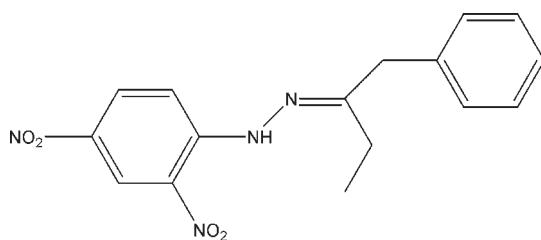
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Key indicators: single-crystal X-ray study; $T = 120\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$; R factor = 0.041; wR factor = 0.101; data-to-parameter ratio = 16.2.

In the title compound, $\text{C}_{16}\text{H}_{16}\text{N}_4\text{O}_4$, the dihedral angle between the aromatic rings is $79.04(8)^\circ$ and an intramolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bond occurs. In the crystal, weak $\text{C}-\text{H}\cdot\cdot\text{O}$ and $\text{C}-\text{H}\cdots\pi$ interactions link the molecules, forming sheets.

Related literature

For the structure of the related 2,4-dinitrophenyl hydrazine, see: Wardell *et al.* (2006). For graph-set notation, see: Bernstein *et al.* (1995).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{16}\text{N}_4\text{O}_4$
 $M_r = 328.33$
Monoclinic, $P2_1/n$

$a = 15.8919(13)\text{ \AA}$
 $b = 4.9446(3)\text{ \AA}$
 $c = 20.7397(17)\text{ \AA}$

$\beta = 105.267(5)^\circ$
 $V = 1572.2(2)\text{ \AA}^3$
 $Z = 4$
Mo $K\alpha$ radiation

$\mu = 0.10\text{ mm}^{-1}$
 $T = 120\text{ K}$
 $0.30 \times 0.05 \times 0.02\text{ mm}$

Data collection

Bruker SMART APEXII
diffractometer
Absorption correction: multi-scan
(*SADABS*; Bruker, 2004)
 $T_{\min} = 0.970$, $T_{\max} = 0.998$

14305 measured reflections
3531 independent reflections
2418 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.101$
 $S = 1.01$
3531 reflections
218 parameters

1 restraint
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.25\text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.19\text{ e \AA}^{-3}$

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1 \cdots O122	0.85	1.95	2.5966 (17)	132
C3—H3A \cdots O142 ⁱ	0.99	2.50	3.432 (2)	158
C32—H32 \cdots O142 ⁱ	0.95	2.52	3.349 (2)	146
C3—H3B \cdots Cg2 ⁱⁱ	0.99	2.75	3.534 (2)	136

Symmetry codes: (i) $x + \frac{1}{2}, -y + \frac{1}{2}, z + \frac{1}{2}$; (ii) $x, y - 1, z$. Cg2 is the centroid of C31–C36 ring.

Data collection: *APEX2* (Bruker, 2004); cell refinement: *APEX2* and *SAINT* (Bruker, 2004); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEPII* (Johnson, 1976) and *PLATON* (Spek, 2009); software used to prepare material for publication: *SHELXL97*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB5130).

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supporting information

Acta Cryst. (2009). E65, o2729 [https://doi.org/10.1107/S1600536809041178]

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S1. Comment

The molecular geometry and conformation is as expected taking account of electronic repulsions and steric effects. The orange colour is caused by the conjugation of the nitrophenyl with the $-N=N-$ group. The backbone of the molecule is essentially planar with only the methyl group, C21, and the phenyl group attached to C3 lying out of the plane of the molecule, Fig. 1.

Atom N1 forms an intramolecular hydrogen bond *via* H1 with atom O122 so forming an $R(6)$ ring, (Bernstein *et al.*, 1995). Atom O142 acts as a hydrogen bond acceptor from donor atoms C3, *via* H3B, and C32, *via* H32, thus forming an $R^2_1(5)$ ring which links screw-related molecules into chains running parallel to (101). These chains are linked to form sheets running parallel to the *b* axis by the C—H $\cdots\pi$ contact in which C3 acts as a donor atom *via* H3B to the phenyl ring containing C31, Fig. 2.

There is a short intermolecular nitro to nitro group contact between O122 \cdots N12($-x + 1.5, y + 1/2, -z + 1/2$) of 2.76 Å.

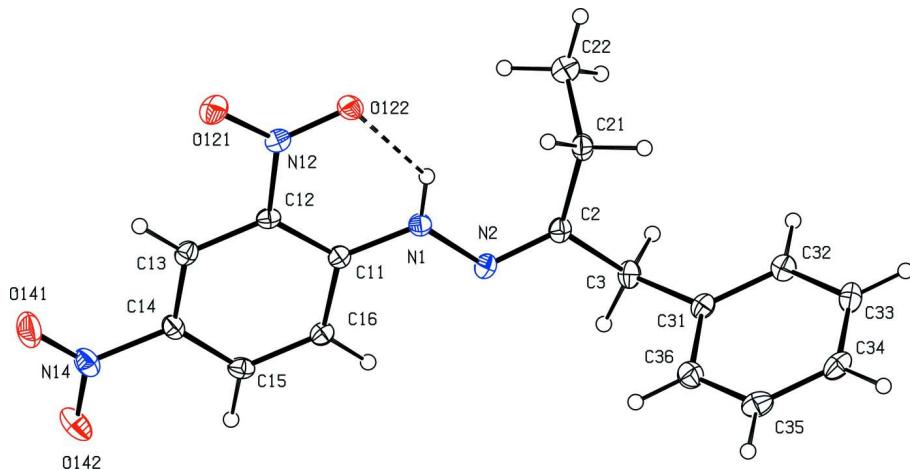
The relevant bonds and angles compare well with those in 2,4-dinitrophenylhydrazine: Wardell *et al.* 2006.

S2. Experimental

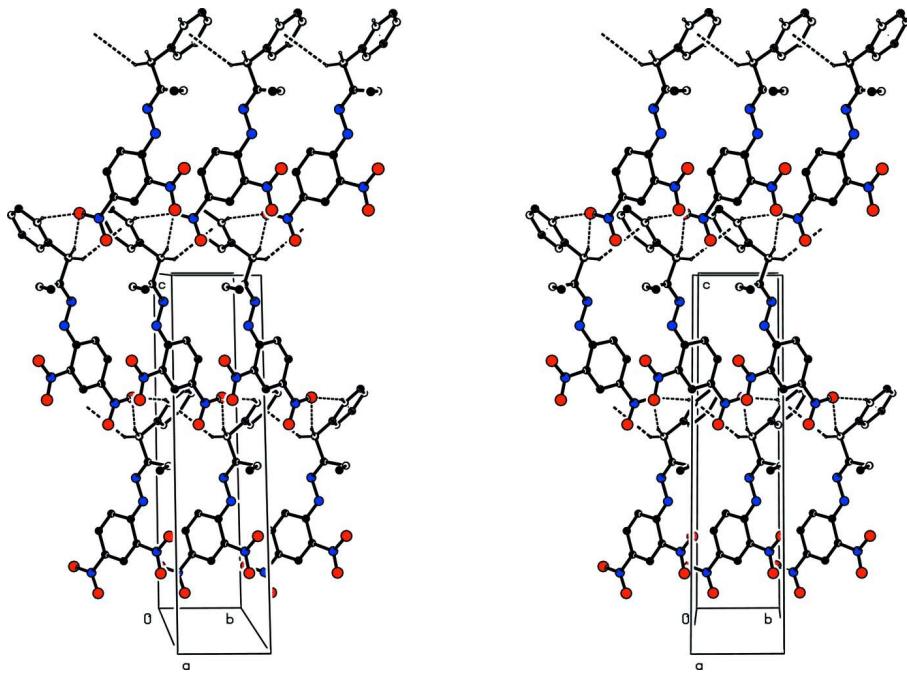
The title compound was obtained from the condensation reaction of 1-phenyl-2-butanone with dinitrophenylhydrazine. The liquid 1-phenyl-2-butanone (2.0 mmol) was added to a warm solution (323 K) of 2,4-dinitrophenylhydrazine (2.5 mmol) in a mixture of 10 ml of ethanol/ 1 ml of HCl (37%) and allowed to react for 30 minutes. The resulting mixture was extracted with ethylacetate. A solid product was obtained after evaporation of the solvent this was first re-crystallized with ethanol and then with ethylacetate. Slow evaporation of a dichloromethane solution gave orange needles of (I).

S3. Refinement

H atoms were treated as riding atoms with C—H(aromatic), 0.95 Å, C—H2(aliphatic), 0.99 Å, C—H(methyl), 0.98 Å. The H atom attached to N1 was located on a difference map, fixed to 0.85 Å, and then refined as a riding atom. The reflections 101 and $\bar{1}01$ were omitted from the refinement as they were obscured by the beamstop.

**Figure 1**

A view of (I) with displacement ellipsoids drawn at the 30% probability level.

**Figure 2**

A stereoview of part of the crystal structure of (I), showing part of the sheet formed by the C—H..O and C—H.. π interactions. Hydrogen atoms not involved in the motifs are not included.

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Crystal data

C₁₆H₁₄N₄O₄

M_r = 328.33

Monoclinic, P2₁/n

Hall symbol: -P 2yn

a = 15.8919 (13) Å

b = 4.9446 (3) Å

c = 20.7397 (17) Å

β = 105.267 (5) $^\circ$

V = 1572.2 (2) Å³

Z = 4

F(000) = 688

D_x = 1.387 Mg m⁻³

Melting point: not measured K
 Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$
 Cell parameters from 2030 reflections
 $\theta = 3.1\text{--}26.5^\circ$

$\mu = 0.10 \text{ mm}^{-1}$
 $T = 120 \text{ K}$
 Needle, orange
 $0.30 \times 0.05 \times 0.02 \text{ mm}$

Data collection

Bruker SMART APEX
 diffractometer
 Radiation source: fine-focus sealed tube
 Graphite monochromator
 Detector resolution: 8.333 pixels mm^{-1}
 ω scans
 Absorption correction: multi-scan
 (*SADABS*; Bruker, 2004)
 $T_{\min} = 0.970$, $T_{\max} = 0.998$

14305 measured reflections
 3531 independent reflections
 2418 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$
 $\theta_{\max} = 27.3^\circ$, $\theta_{\min} = 1.5^\circ$
 $h = -20 \rightarrow 20$
 $k = -5 \rightarrow 6$
 $l = -26 \rightarrow 26$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.041$
 $wR(F^2) = 0.101$
 $S = 1.01$
 3531 reflections
 218 parameters
 1 restraint
 Primary atom site location: structure-invariant
 direct methods
 Secondary atom site location: difference Fourier
 map

Hydrogen site location: inferred from
 neighbouring sites
 H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.037P)^2 + 0.4505P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.25 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e \AA}^{-3}$
 Extinction correction: *SHELXL97* (Sheldrick,
 2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
 Extinction coefficient: 0.0044 (11)

Special details

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

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

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

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O121	0.65610 (8)	0.8996 (2)	0.17034 (6)	0.0325 (3)
O122	0.70928 (7)	0.9899 (2)	0.27500 (6)	0.0276 (3)
O141	0.44704 (9)	0.2157 (3)	0.09414 (6)	0.0454 (4)
O142	0.41347 (8)	-0.0583 (3)	0.16477 (7)	0.0427 (4)
N1	0.66460 (8)	0.7090 (3)	0.36646 (6)	0.0225 (3)
H1	0.6995	0.8261	0.3582	0.027*
N2	0.66485 (9)	0.6313 (3)	0.43054 (6)	0.0225 (3)
N12	0.66156 (9)	0.8554 (3)	0.22955 (7)	0.0225 (3)
N14	0.45198 (9)	0.1408 (3)	0.15136 (7)	0.0301 (3)
C2	0.71757 (10)	0.7597 (3)	0.47833 (8)	0.0220 (3)

C3	0.71901 (11)	0.6653 (3)	0.54758 (8)	0.0250 (4)
H3A	0.7789	0.6066	0.5708	0.030*
H3B	0.6800	0.5068	0.5442	0.030*
C11	0.61365 (10)	0.5756 (3)	0.31389 (7)	0.0195 (3)
C12	0.61004 (10)	0.6396 (3)	0.24669 (8)	0.0194 (3)
C13	0.55682 (10)	0.4984 (3)	0.19381 (8)	0.0228 (4)
H13	0.5557	0.5436	0.1491	0.027*
C14	0.50597 (10)	0.2934 (3)	0.20667 (8)	0.0227 (4)
C15	0.50676 (10)	0.2231 (3)	0.27206 (8)	0.0235 (4)
H15	0.4710	0.0798	0.2801	0.028*
C16	0.55915 (10)	0.3614 (3)	0.32419 (8)	0.0223 (4)
H16	0.5592	0.3133	0.3686	0.027*
C21	0.77979 (11)	0.9796 (3)	0.47154 (8)	0.0269 (4)
H21A	0.7905	1.0997	0.5111	0.032*
H21B	0.7532	1.0896	0.4315	0.032*
C22	0.86642 (12)	0.8638 (4)	0.46542 (10)	0.0410 (5)
H22A	0.9053	1.0120	0.4607	0.061*
H22B	0.8560	0.7461	0.4261	0.061*
H22C	0.8936	0.7587	0.5056	0.061*
C31	0.69053 (11)	0.8820 (3)	0.58860 (8)	0.0228 (4)
C32	0.74933 (11)	1.0031 (3)	0.64209 (8)	0.0257 (4)
H32	0.8088	0.9494	0.6534	0.031*
C33	0.72226 (12)	1.2021 (3)	0.67932 (8)	0.0300 (4)
H33	0.7633	1.2831	0.7160	0.036*
C34	0.63628 (12)	1.2830 (4)	0.66341 (9)	0.0317 (4)
H34	0.6179	1.4195	0.6890	0.038*
C35	0.57701 (12)	1.1645 (4)	0.61015 (9)	0.0332 (4)
H35	0.5177	1.2198	0.5989	0.040*
C36	0.60383 (11)	0.9647 (4)	0.57296 (9)	0.0300 (4)
H36	0.5626	0.8836	0.5365	0.036*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O121	0.0417 (8)	0.0314 (7)	0.0260 (6)	-0.0003 (6)	0.0118 (6)	0.0090 (5)
O122	0.0283 (7)	0.0230 (6)	0.0326 (7)	-0.0063 (5)	0.0100 (5)	-0.0043 (5)
O141	0.0493 (9)	0.0554 (9)	0.0250 (7)	-0.0098 (7)	-0.0019 (6)	-0.0049 (6)
O142	0.0380 (8)	0.0326 (7)	0.0493 (8)	-0.0125 (6)	-0.0030 (6)	-0.0037 (6)
N1	0.0242 (7)	0.0225 (7)	0.0210 (7)	-0.0041 (6)	0.0067 (6)	0.0003 (6)
N2	0.0256 (7)	0.0230 (7)	0.0191 (7)	0.0027 (6)	0.0063 (6)	0.0008 (6)
N12	0.0228 (7)	0.0203 (7)	0.0256 (7)	0.0040 (6)	0.0086 (6)	0.0033 (6)
N14	0.0232 (8)	0.0291 (8)	0.0331 (8)	0.0018 (6)	-0.0011 (6)	-0.0049 (7)
C2	0.0230 (8)	0.0197 (8)	0.0231 (8)	0.0056 (7)	0.0058 (7)	-0.0023 (7)
C3	0.0298 (9)	0.0224 (8)	0.0212 (8)	0.0029 (7)	0.0040 (7)	0.0004 (7)
C11	0.0188 (8)	0.0171 (8)	0.0226 (8)	0.0040 (6)	0.0056 (6)	-0.0013 (6)
C12	0.0184 (8)	0.0167 (8)	0.0246 (8)	0.0021 (6)	0.0082 (6)	0.0017 (7)
C13	0.0227 (9)	0.0248 (9)	0.0213 (8)	0.0060 (7)	0.0066 (7)	0.0008 (7)
C14	0.0187 (8)	0.0210 (8)	0.0267 (8)	0.0020 (7)	0.0028 (7)	-0.0057 (7)

C15	0.0205 (8)	0.0188 (8)	0.0327 (9)	-0.0007 (7)	0.0097 (7)	-0.0003 (7)
C16	0.0237 (9)	0.0215 (8)	0.0230 (8)	0.0015 (7)	0.0085 (7)	0.0006 (7)
C21	0.0311 (10)	0.0275 (9)	0.0217 (8)	-0.0036 (7)	0.0063 (7)	-0.0053 (7)
C22	0.0344 (11)	0.0536 (13)	0.0383 (10)	-0.0071 (9)	0.0157 (9)	-0.0107 (10)
C31	0.0286 (9)	0.0205 (8)	0.0204 (8)	0.0025 (7)	0.0084 (7)	0.0044 (7)
C32	0.0272 (9)	0.0245 (9)	0.0246 (8)	0.0035 (7)	0.0056 (7)	0.0028 (7)
C33	0.0404 (11)	0.0275 (9)	0.0219 (8)	-0.0012 (8)	0.0081 (8)	-0.0024 (7)
C34	0.0413 (11)	0.0275 (10)	0.0323 (9)	0.0049 (8)	0.0206 (8)	-0.0001 (8)
C35	0.0288 (10)	0.0361 (10)	0.0382 (10)	0.0070 (8)	0.0149 (8)	0.0046 (9)
C36	0.0282 (10)	0.0329 (10)	0.0278 (9)	0.0005 (8)	0.0053 (7)	-0.0002 (8)

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

O121—N12	1.2276 (16)	C15—C16	1.362 (2)
O122—N12	1.2370 (17)	C15—H15	0.9500
O141—N14	1.2258 (18)	C16—H16	0.9500
O142—N14	1.2292 (19)	C21—C22	1.527 (2)
N1—C11	1.3469 (19)	C21—H21A	0.9900
N1—N2	1.3823 (17)	C21—H21B	0.9900
N1—H1	0.85	C22—H22A	0.9800
N2—C2	1.284 (2)	C22—H22B	0.9800
N12—C12	1.445 (2)	C22—H22C	0.9800
N14—C14	1.450 (2)	C31—C32	1.385 (2)
C2—C21	1.501 (2)	C31—C36	1.391 (2)
C2—C3	1.505 (2)	C32—C33	1.387 (2)
C3—C31	1.510 (2)	C32—H32	0.9500
C3—H3A	0.9900	C33—C34	1.378 (2)
C3—H3B	0.9900	C33—H33	0.9500
C11—C12	1.416 (2)	C34—C35	1.379 (3)
C11—C16	1.420 (2)	C34—H34	0.9500
C12—C13	1.385 (2)	C35—C36	1.388 (2)
C13—C14	1.366 (2)	C35—H35	0.9500
C13—H13	0.9500	C36—H36	0.9500
C14—C15	1.397 (2)		
C11—N1—N2	119.42 (13)	C15—C16—C11	121.65 (15)
C11—N1—H1	117.1	C15—C16—H16	119.2
N2—N1—H1	123.1	C11—C16—H16	119.2
C2—N2—N1	116.15 (13)	C2—C21—C22	111.49 (14)
O121—N12—O122	122.28 (13)	C2—C21—H21A	109.3
O121—N12—C12	118.84 (13)	C22—C21—H21A	109.3
O122—N12—C12	118.88 (13)	C2—C21—H21B	109.3
O141—N14—O142	123.53 (15)	C22—C21—H21B	109.3
O141—N14—C14	118.82 (15)	H21A—C21—H21B	108.0
O142—N14—C14	117.64 (15)	C21—C22—H22A	109.5
N2—C2—C21	126.70 (14)	C21—C22—H22B	109.5
N2—C2—C3	115.22 (14)	H22A—C22—H22B	109.5
C21—C2—C3	117.99 (14)	C21—C22—H22C	109.5

C2—C3—C31	112.72 (13)	H22A—C22—H22C	109.5
C2—C3—H3A	109.0	H22B—C22—H22C	109.5
C31—C3—H3A	109.0	C32—C31—C36	118.53 (15)
C2—C3—H3B	109.0	C32—C31—C3	121.28 (15)
C31—C3—H3B	109.0	C36—C31—C3	120.19 (15)
H3A—C3—H3B	107.8	C31—C32—C33	120.67 (16)
N1—C11—C12	123.19 (14)	C31—C32—H32	119.7
N1—C11—C16	120.26 (14)	C33—C32—H32	119.7
C12—C11—C16	116.54 (14)	C34—C33—C32	120.39 (17)
C13—C12—C11	121.67 (14)	C34—C33—H33	119.8
C13—C12—N12	116.42 (14)	C32—C33—H33	119.8
C11—C12—N12	121.91 (14)	C33—C34—C35	119.58 (16)
C14—C13—C12	119.28 (15)	C33—C34—H34	120.2
C14—C13—H13	120.4	C35—C34—H34	120.2
C12—C13—H13	120.4	C34—C35—C36	120.17 (17)
C13—C14—C15	121.32 (15)	C34—C35—H35	119.9
C13—C14—N14	119.28 (15)	C36—C35—H35	119.9
C15—C14—N14	119.38 (15)	C35—C36—C31	120.67 (17)
C16—C15—C14	119.54 (15)	C35—C36—H36	119.7
C16—C15—H15	120.2	C31—C36—H36	119.7
C14—C15—H15	120.2		
C11—N1—N2—C2	177.22 (14)	O142—N14—C14—C13	173.19 (15)
N1—N2—C2—C21	-1.4 (2)	O141—N14—C14—C15	174.98 (15)
N1—N2—C2—C3	-177.96 (12)	O142—N14—C14—C15	-5.0 (2)
N2—C2—C3—C31	-117.81 (16)	C13—C14—C15—C16	0.1 (2)
C21—C2—C3—C31	65.32 (19)	N14—C14—C15—C16	178.26 (14)
N2—N1—C11—C12	-179.33 (13)	C14—C15—C16—C11	-0.3 (2)
N2—N1—C11—C16	1.5 (2)	N1—C11—C16—C15	179.86 (14)
N1—C11—C12—C13	-179.94 (14)	C12—C11—C16—C15	0.6 (2)
C16—C11—C12—C13	-0.7 (2)	N2—C2—C21—C22	-86.2 (2)
N1—C11—C12—N12	0.1 (2)	C3—C2—C21—C22	90.25 (17)
C16—C11—C12—N12	179.29 (13)	C2—C3—C31—C32	-109.84 (17)
O121—N12—C12—C13	-0.6 (2)	C2—C3—C31—C36	70.0 (2)
O122—N12—C12—C13	179.04 (13)	C36—C31—C32—C33	0.1 (2)
O121—N12—C12—C11	179.42 (14)	C3—C31—C32—C33	179.95 (15)
O122—N12—C12—C11	-1.0 (2)	C31—C32—C33—C34	-0.2 (3)
C11—C12—C13—C14	0.5 (2)	C32—C33—C34—C35	0.1 (3)
N12—C12—C13—C14	-179.50 (13)	C33—C34—C35—C36	0.2 (3)
C12—C13—C14—C15	-0.2 (2)	C34—C35—C36—C31	-0.2 (3)
C12—C13—C14—N14	-178.37 (14)	C32—C31—C36—C35	0.1 (2)
O141—N14—C14—C13	-6.8 (2)	C3—C31—C36—C35	-179.73 (15)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O122	0.85	1.95	2.5966 (17)	132
C3—H3A···O142 ⁱ	0.99	2.50	3.432 (2)	158

C32—H32···O142 ⁱ	0.95	2.52	3.349 (2)	146
C3—H3B···Cg2 ⁱⁱ	0.99	2.75	3.534 (2)	136

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