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***N*-(2-Oxo-2*H*-chromen-3-yl)benzamide**Mukesh M. Jotani,^{a‡} Bharat B. Baldaniya^b and Edward R. T. Tiekink^{c*}

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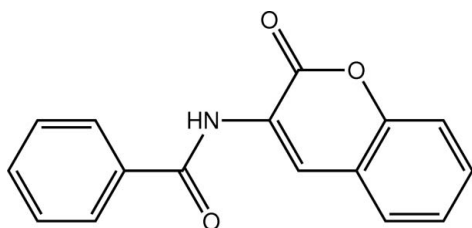
Received 2 March 2010; accepted 4 March 2010

Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.040; wR factor = 0.118; data-to-parameter ratio = 15.4.

The phenyl ring in title molecule, $\text{C}_{16}\text{H}_{11}\text{NO}_3$, forms a dihedral angle of 7.69 (6)° with the fused ring system. The observed conformation is stabilized by intramolecular $\text{N}-\text{H}\cdots\text{O}$ and $\text{C}-\text{H}\cdots\text{O}$ interactions. In the crystal, supramolecular chains are formed along the b axis which are mediated by $\pi-\pi$ interactions [centroid-centroid distance = 3.614 (2) Å].

Related literature

For the biological activity of imidazoles, see: Yohjiro *et al.* (1990). For the anti-inflammatory activity of the title compound, see: Maddi *et al.* (2007). Semi-empirical quantum chemical calculations were performed using *MOPAC2009* Stewart (2009).



Experimental

Crystal data

$\text{C}_{16}\text{H}_{11}\text{NO}_3$
 $M_r = 265.26$
Monoclinic, $P2_1/c$
 $a = 12.519$ (4) Å
 $b = 4.748$ (3) Å

$c = 21.167$ (4) Å
 $\beta = 102.044$ (3)°
 $V = 1230.5$ (9) Å³
 $Z = 4$
Mo $K\alpha$ radiation

$\mu = 0.10$ mm⁻¹
 $T = 293$ K

 $0.40 \times 0.22 \times 0.15$ mm

Data collection

Bruker SMART APEX CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.945$, $T_{\max} = 0.995$

13295 measured reflections
2827 independent reflections
2029 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.118$
 $S = 1.11$
2827 reflections
184 parameters
1 restraint

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.18$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.18$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|---------------------------------------|----------|-------------|-------------|---------------|
| $\text{N1}-\text{H1n}\cdots\text{O1}$ | 0.87 (1) | 2.24 (2) | 2.659 (2) | 110 (1) |
| $\text{C8}-\text{H8}\cdots\text{O3}$ | 0.93 | 2.24 | 2.822 (3) | 120 |

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: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *DIAMOND* (Brandenburg, 2006); software used to prepare material for publication: *publCIF* (Westrip, 2010).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HG2654).

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supporting information

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N*-(2-Oxo-2*H*-chromen-3-yl)benzamide*Mukesh M. Jotani, Bharat B. Baldaniya and Edward R. T. Tiekink****S1. Comment**

Oxazoles are very useful synthetic intermediates used for the generation of imidazoles that possess a wide spectrum of biological activities such as herbicidal, anti-bacterial, anti-fungal, *etc.* (Yohjiro *et al.*, 1990). During attempts designed to synthesize oxazoles containing various substituents at different positions in the benzene ring, the title compound, (I), was obtained unexpectedly by the formation of a chromen derivative instead of the anticipated oxazole. Compound (I) is a known species and has been shown to be pharmaceutically important as it possesses anti-inflammatory activity (Maddi *et al.*, 2007).

There is a twist in the molecule of (I), Fig. 1, so that the pendent benzene ring is not co-planar with the rest of the molecule. This is seen in the value of the O3–C10–C11–C12 torsion angle of $-166.43(15)^\circ$, and in the dihedral formed between the fused ring system and benzene ring of $7.69(6)^\circ$. The overall conformation of the molecule is stabilised by intramolecular N–H \cdots O and C–H \cdots O interactions, Table 1, which close S(5) and S(6) hydrogen bond ring motifs, respectively. The most prominent feature of the crystal packing is the formation of supramolecular chains along the *b* axis mediated by π – π interactions between the ring centroids of the (O2,C1,C2,C7–C9) and (C2–C7)^{*i*} rings of $3.614(2) \text{ \AA}$ of translationally related molecules; symmetry operation *i*: $x, 1+y, z$.

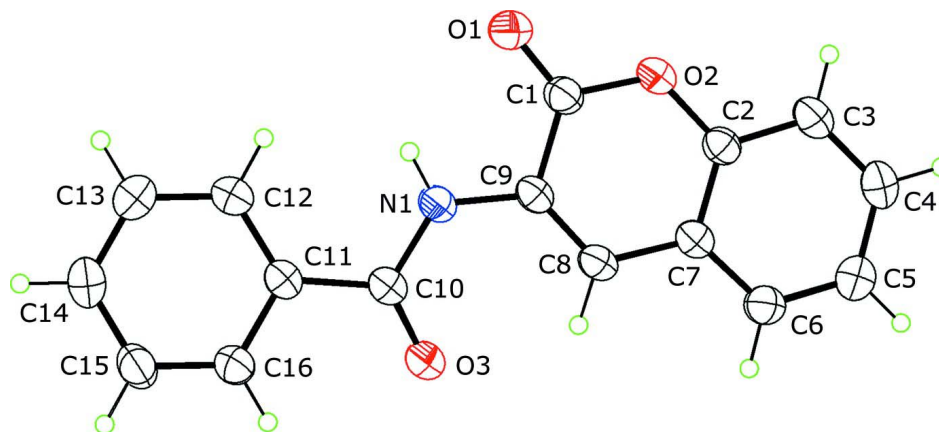
Semi-empirical Quantum Chemical Calculations were performed on experimental structure using MOPAC2009 program (Stewart, 2009) to optimize the molecule with the Austin Model 1 (AM1) approximation, together with the restricted Hartree-Fock closed-shell wavefunction. Minimisations were terminated at an r.m.s. gradient of less than $1.0 \text{ kJ mol}^{-1} \text{ \AA}^{-1}$. These calculations gave a heat of formation = -213.437 kJ for (I). The ionization potential, dipole moment and self consistency field (SCF) factor were calculated as 9.033 eV , 1.798 Debye and 73 , respectively.

S2. Experimental

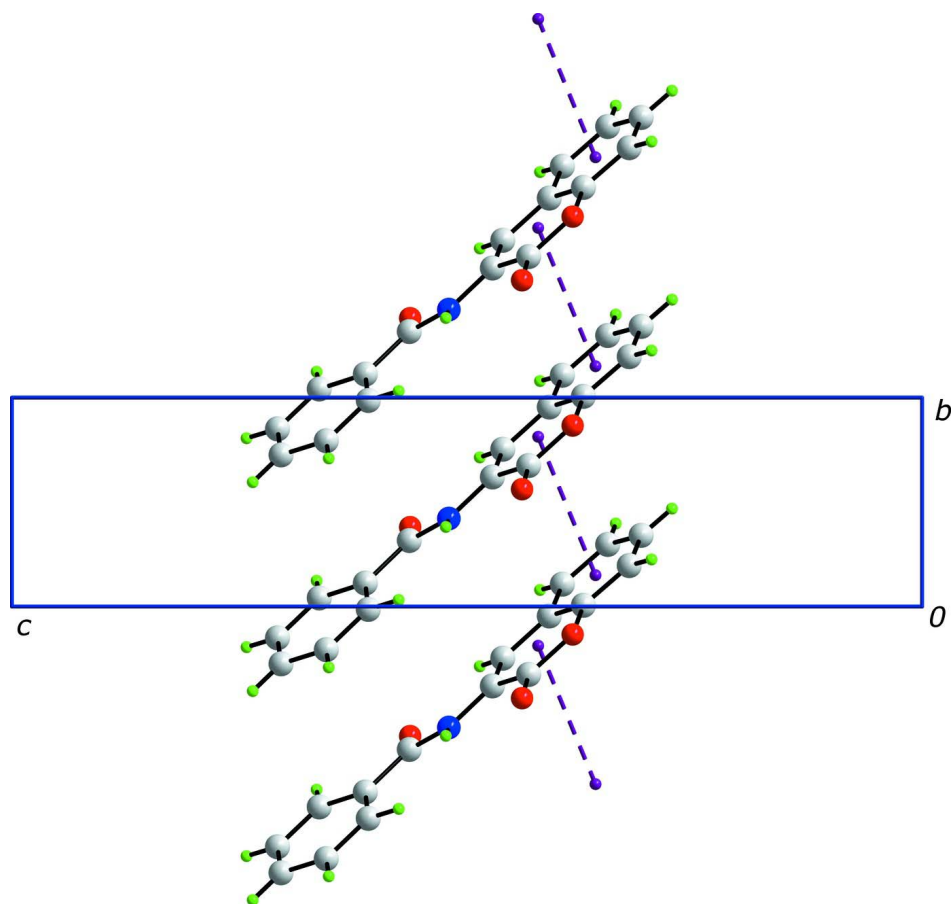
A mixture of orthohydroxy benzaldehyde (0.25 mol), benzoyl amino acetic acid (0.25 mol), acetyl acetate (0.30 mol), and anhydrous sodium acetate (0.25 mol) were taken in a 500 ml round bottom flask and heated on an electric hot plate with constant stirring. After complete liquefaction, the flask was transferred to a sand bath and further heated for 2 h. Ethanol (100 ml) was added slowly to the flask and the mixture was allowed to stand overnight. The crystalline product obtained was filtered with ice-cold alcohol and then with boiling water. The crude product was crystallised from ethanol (95%) to obtain the final product (75 % yield, m.pt. 432 K). The colorless crystals were obtained by slow evaporation from an ethanol solution.

S3. Refinement

The C-bound H atoms were geometrically placed (C–H = 0.93 \AA) and refined as riding with $U_{iso}(\text{H}) = 1.2U_{eq}(\text{parent atom})$. The position of the N–H atom was refined with $U_{iso}(\text{H}) = 1.2U_{eq}(\text{N})$.

**Figure 1**

The molecular structure of (I) showing the atom-labelling scheme and displacement ellipsoids at the 35% probability level.

**Figure 2**

A supramolecular chain aligned along the *b* axis in (I), mediated by π - π interactions (purple dashed lines). Colour code: O, red; N, blue; C, grey; and H, green.

N*-(2-Oxo-2*H*-chromen-3-yl)benzamideCrystal data*C₁₆H₁₁NO₃ $M_r = 265.26$ Monoclinic, $P2_1/c$ Hall symbol: - P 2ybc $a = 12.519$ (4) Å $b = 4.748$ (3) Å $c = 21.167$ (4) Å $\beta = 102.044$ (3)° $V = 1230.5$ (9) Å³ $Z = 4$ $F(000) = 552$ $D_x = 1.432$ Mg m⁻³Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 3699 reflections

 $\theta = 2.3$ – 29.6 ° $\mu = 0.10$ mm⁻¹ $T = 293$ K

Block, colourless

 $0.40 \times 0.22 \times 0.15$ mm*Data collection*

Bruker SMART APEX CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

 ω and φ scans

Absorption correction: multi-scan

(SADABS; Sheldrick, 1996)

 $T_{\min} = 0.945$, $T_{\max} = 0.995$

13295 measured reflections

2827 independent reflections

2029 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.025$ $\theta_{\text{max}} = 27.5$ °, $\theta_{\text{min}} = 1.7$ ° $h = -16$ → 15 $k = -6$ → 5 $l = -27$ → 27 *Refinement*Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.040$ $wR(F^2) = 0.118$ $S = 1.11$

2827 reflections

184 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 atoms treated by a mixture of independent and constrained refinement

 $w = 1/[\sigma^2(F_o^2) + (0.0491P)^2 + 0.2124P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.18$ e Å⁻³ $\Delta\rho_{\text{min}} = -0.18$ e Å⁻³*Special details*

Geometry. All s.u.'s (except the s.u. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell s.u.'s are taken into account individually in the estimation of s.u.'s in distances, angles and torsion angles; correlations between s.u.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell s.u.'s is used for estimating s.u.'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 $F^2 > 2\sigma(F^2)$ 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 (Å²)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|----|--------------|------------|-------------|----------------------------------|
| O1 | 0.57658 (10) | 0.4385 (3) | 0.56046 (6) | 0.0705 (4) |
| O2 | 0.69047 (9) | 0.1338 (2) | 0.61603 (5) | 0.0526 (3) |
| O3 | 0.85472 (10) | 0.6200 (3) | 0.43753 (6) | 0.0667 (4) |

| | | | | |
|-----|--------------|-------------|-------------|------------|
| N1 | 0.70508 (10) | 0.5787 (3) | 0.48003 (6) | 0.0443 (3) |
| H1N | 0.6366 (8) | 0.619 (4) | 0.4767 (8) | 0.053* |
| C1 | 0.66406 (13) | 0.3252 (3) | 0.56770 (7) | 0.0469 (4) |
| C2 | 0.78915 (12) | -0.0059 (3) | 0.62787 (7) | 0.0426 (3) |
| C3 | 0.80701 (14) | -0.1960 (3) | 0.67819 (7) | 0.0534 (4) |
| H3 | 0.7545 | -0.2255 | 0.7027 | 0.064* |
| C4 | 0.90379 (15) | -0.3405 (4) | 0.69126 (8) | 0.0555 (4) |
| H4 | 0.9174 | -0.4688 | 0.7252 | 0.067* |
| C5 | 0.98102 (14) | -0.2972 (4) | 0.65453 (8) | 0.0552 (4) |
| H5 | 1.0463 | -0.3974 | 0.6635 | 0.066* |
| C6 | 0.96208 (13) | -0.1070 (3) | 0.60475 (8) | 0.0509 (4) |
| H6 | 1.0148 | -0.0793 | 0.5802 | 0.061* |
| C7 | 0.86511 (12) | 0.0452 (3) | 0.59038 (7) | 0.0414 (3) |
| C8 | 0.84018 (12) | 0.2473 (3) | 0.53927 (7) | 0.0440 (4) |
| H8 | 0.8909 | 0.2863 | 0.5140 | 0.053* |
| C9 | 0.74390 (12) | 0.3800 (3) | 0.52784 (6) | 0.0403 (3) |
| C10 | 0.76069 (13) | 0.6840 (3) | 0.43656 (7) | 0.0440 (4) |
| C11 | 0.70075 (12) | 0.8878 (3) | 0.38827 (7) | 0.0408 (3) |
| C12 | 0.60296 (13) | 1.0138 (3) | 0.39149 (7) | 0.0500 (4) |
| H12 | 0.5684 | 0.9685 | 0.4250 | 0.060* |
| C13 | 0.55600 (14) | 1.2065 (4) | 0.34538 (8) | 0.0579 (4) |
| H13 | 0.4901 | 1.2911 | 0.3481 | 0.069* |
| C14 | 0.60574 (15) | 1.2743 (4) | 0.29565 (8) | 0.0584 (5) |
| H14 | 0.5738 | 1.4047 | 0.2646 | 0.070* |
| C15 | 0.70244 (15) | 1.1497 (4) | 0.29178 (8) | 0.0590 (5) |
| H15 | 0.7360 | 1.1945 | 0.2578 | 0.071* |
| C16 | 0.75030 (14) | 0.9590 (3) | 0.33763 (7) | 0.0510 (4) |
| H16 | 0.8165 | 0.8765 | 0.3348 | 0.061* |

Atomic displacement parameters (Å²)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|-------------|------------|-------------|
| O1 | 0.0528 (7) | 0.0950 (10) | 0.0687 (8) | 0.0157 (7) | 0.0238 (6) | 0.0260 (7) |
| O2 | 0.0523 (7) | 0.0612 (7) | 0.0488 (6) | 0.0031 (5) | 0.0209 (5) | 0.0136 (5) |
| O3 | 0.0563 (7) | 0.0819 (9) | 0.0682 (8) | 0.0137 (6) | 0.0275 (6) | 0.0313 (7) |
| N1 | 0.0458 (7) | 0.0477 (7) | 0.0413 (7) | -0.0018 (6) | 0.0135 (6) | 0.0049 (5) |
| C1 | 0.0461 (9) | 0.0543 (9) | 0.0415 (8) | -0.0016 (7) | 0.0118 (6) | 0.0042 (7) |
| C2 | 0.0472 (8) | 0.0423 (8) | 0.0390 (7) | -0.0038 (6) | 0.0106 (6) | -0.0016 (6) |
| C3 | 0.0634 (10) | 0.0557 (9) | 0.0434 (8) | -0.0038 (8) | 0.0167 (7) | 0.0066 (7) |
| C4 | 0.0685 (11) | 0.0486 (9) | 0.0471 (9) | -0.0004 (8) | 0.0063 (8) | 0.0074 (7) |
| C5 | 0.0551 (10) | 0.0509 (9) | 0.0572 (10) | 0.0028 (8) | 0.0064 (8) | 0.0012 (8) |
| C6 | 0.0508 (9) | 0.0525 (9) | 0.0512 (9) | -0.0005 (7) | 0.0149 (7) | -0.0009 (7) |
| C7 | 0.0491 (8) | 0.0388 (7) | 0.0373 (7) | -0.0063 (6) | 0.0112 (6) | -0.0043 (6) |
| C8 | 0.0504 (9) | 0.0449 (8) | 0.0401 (7) | -0.0046 (7) | 0.0175 (6) | -0.0009 (6) |
| C9 | 0.0478 (8) | 0.0399 (7) | 0.0344 (7) | -0.0063 (6) | 0.0115 (6) | -0.0014 (6) |
| C10 | 0.0487 (9) | 0.0443 (8) | 0.0407 (8) | -0.0043 (7) | 0.0130 (6) | 0.0005 (6) |
| C11 | 0.0474 (8) | 0.0375 (7) | 0.0376 (7) | -0.0066 (6) | 0.0090 (6) | -0.0024 (6) |
| C12 | 0.0514 (9) | 0.0547 (9) | 0.0462 (8) | -0.0042 (8) | 0.0157 (7) | 0.0005 (7) |

| | | | | | | |
|-----|-------------|-------------|-------------|------------|------------|-------------|
| C13 | 0.0512 (10) | 0.0578 (10) | 0.0628 (10) | 0.0063 (8) | 0.0075 (8) | -0.0003 (8) |
| C14 | 0.0706 (12) | 0.0501 (9) | 0.0506 (9) | 0.0028 (8) | 0.0037 (8) | 0.0078 (8) |
| C15 | 0.0740 (12) | 0.0572 (10) | 0.0485 (9) | 0.0024 (9) | 0.0191 (8) | 0.0113 (8) |
| C16 | 0.0570 (9) | 0.0527 (9) | 0.0463 (8) | 0.0035 (8) | 0.0177 (7) | 0.0036 (7) |

Geometric parameters (Å, °)

| | | | |
|------------|-------------|-------------|-------------|
| O1—C1 | 1.2009 (19) | C6—H6 | 0.9300 |
| O2—C1 | 1.3567 (18) | C7—C8 | 1.431 (2) |
| O2—C2 | 1.3783 (18) | C8—C9 | 1.336 (2) |
| O3—C10 | 1.2118 (18) | C8—H8 | 0.9300 |
| N1—C10 | 1.3596 (19) | C10—C11 | 1.492 (2) |
| N1—C9 | 1.3947 (19) | C11—C12 | 1.377 (2) |
| N1—H1N | 0.867 (9) | C11—C16 | 1.388 (2) |
| C1—C9 | 1.460 (2) | C12—C13 | 1.377 (2) |
| C2—C3 | 1.378 (2) | C12—H12 | 0.9300 |
| C2—C7 | 1.381 (2) | C13—C14 | 1.369 (3) |
| C3—C4 | 1.369 (2) | C13—H13 | 0.9300 |
| C3—H3 | 0.9300 | C14—C15 | 1.365 (2) |
| C4—C5 | 1.377 (2) | C14—H14 | 0.9300 |
| C4—H4 | 0.9300 | C15—C16 | 1.371 (2) |
| C5—C6 | 1.370 (2) | C15—H15 | 0.9300 |
| C5—H5 | 0.9300 | C16—H16 | 0.9300 |
| C6—C7 | 1.391 (2) | | |
| C1—O2—C2 | 121.85 (12) | C9—C8—H8 | 120.0 |
| C10—N1—C9 | 126.13 (13) | C7—C8—H8 | 120.0 |
| C10—N1—H1N | 120.0 (11) | C8—C9—N1 | 127.93 (14) |
| C9—N1—H1N | 113.5 (11) | C8—C9—C1 | 120.79 (13) |
| O1—C1—O2 | 117.93 (14) | N1—C9—C1 | 111.28 (13) |
| O1—C1—C9 | 124.25 (14) | O3—C10—N1 | 121.95 (14) |
| O2—C1—C9 | 117.81 (14) | O3—C10—C11 | 121.49 (13) |
| O2—C2—C3 | 116.77 (14) | N1—C10—C11 | 116.55 (14) |
| O2—C2—C7 | 120.63 (13) | C12—C11—C16 | 118.54 (14) |
| C3—C2—C7 | 122.60 (15) | C12—C11—C10 | 124.99 (14) |
| C4—C3—C2 | 118.57 (16) | C16—C11—C10 | 116.44 (14) |
| C4—C3—H3 | 120.7 | C11—C12—C13 | 120.41 (15) |
| C2—C3—H3 | 120.7 | C11—C12—H12 | 119.8 |
| C3—C4—C5 | 120.47 (15) | C13—C12—H12 | 119.8 |
| C3—C4—H4 | 119.8 | C14—C13—C12 | 120.40 (17) |
| C5—C4—H4 | 119.8 | C14—C13—H13 | 119.8 |
| C6—C5—C4 | 120.23 (16) | C12—C13—H13 | 119.8 |
| C6—C5—H5 | 119.9 | C15—C14—C13 | 119.71 (16) |
| C4—C5—H5 | 119.9 | C15—C14—H14 | 120.1 |
| C5—C6—C7 | 120.90 (15) | C13—C14—H14 | 120.1 |
| C5—C6—H6 | 119.5 | C14—C15—C16 | 120.40 (16) |
| C7—C6—H6 | 119.5 | C14—C15—H15 | 119.8 |
| C2—C7—C6 | 117.21 (14) | C16—C15—H15 | 119.8 |

| | | | |
|--------------|--------------|-----------------|--------------|
| C2—C7—C8 | 118.97 (14) | C15—C16—C11 | 120.53 (16) |
| C6—C7—C8 | 123.82 (14) | C15—C16—H16 | 119.7 |
| C9—C8—C7 | 119.94 (14) | C11—C16—H16 | 119.7 |
| C2—O2—C1—O1 | -179.90 (14) | C10—N1—C9—C1 | 177.37 (14) |
| C2—O2—C1—C9 | 0.1 (2) | O1—C1—C9—C8 | 179.34 (16) |
| C1—O2—C2—C3 | -179.91 (13) | O2—C1—C9—C8 | -0.7 (2) |
| C1—O2—C2—C7 | 0.0 (2) | O1—C1—C9—N1 | -0.5 (2) |
| O2—C2—C3—C4 | 179.59 (14) | O2—C1—C9—N1 | 179.48 (12) |
| C7—C2—C3—C4 | -0.3 (2) | C9—N1—C10—O3 | -3.4 (2) |
| C2—C3—C4—C5 | -0.4 (2) | C9—N1—C10—C11 | 177.89 (13) |
| C3—C4—C5—C6 | 0.5 (3) | O3—C10—C11—C12 | -166.43 (15) |
| C4—C5—C6—C7 | 0.0 (2) | N1—C10—C11—C12 | 12.3 (2) |
| O2—C2—C7—C6 | -179.11 (13) | O3—C10—C11—C16 | 11.7 (2) |
| C3—C2—C7—C6 | 0.7 (2) | N1—C10—C11—C16 | -169.62 (14) |
| O2—C2—C7—C8 | 0.5 (2) | C16—C11—C12—C13 | -0.2 (2) |
| C3—C2—C7—C8 | -179.65 (14) | C10—C11—C12—C13 | 177.90 (14) |
| C5—C6—C7—C2 | -0.6 (2) | C11—C12—C13—C14 | 0.3 (3) |
| C5—C6—C7—C8 | 179.82 (14) | C12—C13—C14—C15 | 0.1 (3) |
| C2—C7—C8—C9 | -1.0 (2) | C13—C14—C15—C16 | -0.5 (3) |
| C6—C7—C8—C9 | 178.53 (14) | C14—C15—C16—C11 | 0.6 (3) |
| C7—C8—C9—N1 | -179.05 (13) | C12—C11—C16—C15 | -0.3 (2) |
| C7—C8—C9—C1 | 1.1 (2) | C10—C11—C16—C15 | -178.48 (14) |
| C10—N1—C9—C8 | -2.5 (2) | | |

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—H1n \cdots O1 | 0.87 (1) | 2.24 (2) | 2.659 (2) | 110 (1) |
| C8—H8 \cdots O3 | 0.93 | 2.24 | 2.822 (3) | 120 |