

The cocrystal 2-hydroxy-4-methyl-N-propanoylbenzohydrazide–2-hydroxy-N-(2-hydroxy-4-methylbenzoyl)-6-methylbenzohydrazide (2/1)

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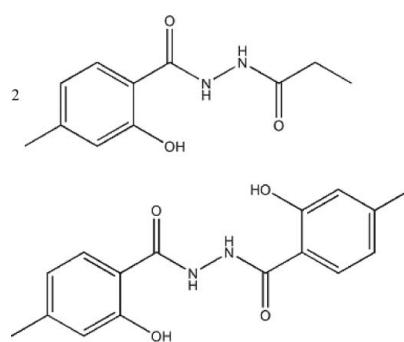
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Key indicators: single-crystal X-ray study; $T = 296\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.003\text{ \AA}$; R factor = 0.037; wR factor = 0.110; data-to-parameter ratio = 16.8.

The asymmetric unit of the title compound, $2\text{C}_{11}\text{H}_{14}\text{N}_2\text{O}_3\cdot\text{C}_{16}\text{H}_{16}\text{N}_2\text{O}_4$, contains one molecule of 2-hydroxy-4-methyl-N-propanoylbenzohydrazide and one-half of a molecule of 2-hydroxy-N-(2-hydroxy-4-methylbenzoyl)-6-methylbenzohydrazide. The latter is located on a centre of inversion. Intramolecular N—H···O interactions stabilize the conformations of both molecules. The crystal structure is stabilized by intermolecular N—H···O and O—H···O hydrogen bonds.

Related literature

For related literature, see: John *et al.* (2007); Majumder *et al.* (2006).



Experimental

Crystal data

$2\text{C}_{11}\text{H}_{14}\text{N}_2\text{O}_3\cdot\text{C}_{16}\text{H}_{16}\text{N}_2\text{O}_4$	$\gamma = 104.448(4)^\circ$
$M_r = 744.79$	$V = 888.8(2)\text{ \AA}^3$
Triclinic, $P\bar{1}$	$Z = 1$
$a = 6.5778(10)\text{ \AA}$	Mo $K\alpha$ radiation
$b = 10.7618(17)\text{ \AA}$	$\mu = 0.10\text{ mm}^{-1}$
$c = 13.936(2)\text{ \AA}$	$T = 296(2)\text{ K}$
$\alpha = 109.522(3)^\circ$	$0.54 \times 0.30 \times 0.25\text{ mm}$
$\beta = 93.608(1)^\circ$	

Data collection

Bruker APEXII diffractometer	5076 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2005)	4125 independent reflections
$T_{\min} = 0.964$, $T_{\max} = 0.975$	1865 reflections with $I > 2\sigma(I)$
	$R_{\text{int}} = 0.022$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.037$	H atoms treated by a mixture of independent and constrained refinement
$wR(F^2) = 0.110$	$\Delta\rho_{\text{max}} = 0.20\text{ e \AA}^{-3}$
$S = 0.90$	$\Delta\rho_{\text{min}} = -0.22\text{ e \AA}^{-3}$
4125 reflections	
246 parameters	

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—H1D···O1	0.86	1.93	2.620 (2)	136
O1—H1E···O4	0.82	1.87	2.682 (2)	169
N2—H2A···O2 ⁱ	0.86	2.03	2.866 (2)	165
N3—H3B···O5	0.84 (2)	1.94 (3)	2.613 (3)	136 (2)
N3—H3B···O4 ⁱⁱ	0.84 (2)	2.37 (3)	2.655 (3)	101 (2)
O5—H5B···O3 ⁱⁱ	0.89 (3)	1.80 (3)	2.685 (2)	175 (2)

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT2793).

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supporting information

Acta Cryst. (2008). E64, o2144 [doi:10.1107/S1600536808033515]

The cocrystal 2-hydroxy-4-methyl-*N*-propanoylbenzohydrazide–2-hydroxy-*N*-(2-hydroxy-4-methylbenzoyl)-6-methylbenzohydrazide (2/1)

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S1. Comment

N-Acylsalicylhydrazides are an interesting class of compounds because of their unique properties. They have been used extensively as ligands in coordination chemistry. *N*-acylsalicylhydrazide compounds show photoluminescence in the solid state by proton transfer from O atom to the imine N atom (Majumder *et al.*, 2006). The nuclearity and the shape of the metallamacrocycles could be modulated by controlling the steric interactions caused by *N*-acyl tails of the ligands (John *et al.*, 2007).

A view of the title structure is illustrated in Fig. 1. The asymmetric unit contains one molecule of 2-hydroxy-4-methyl-*N*-propanoylbenzohydrazide and half a molecule of 2-hydroxy-*N*-(2-hydroxy-4-methylbenzoyl)-6-methylbenzohydrazide.

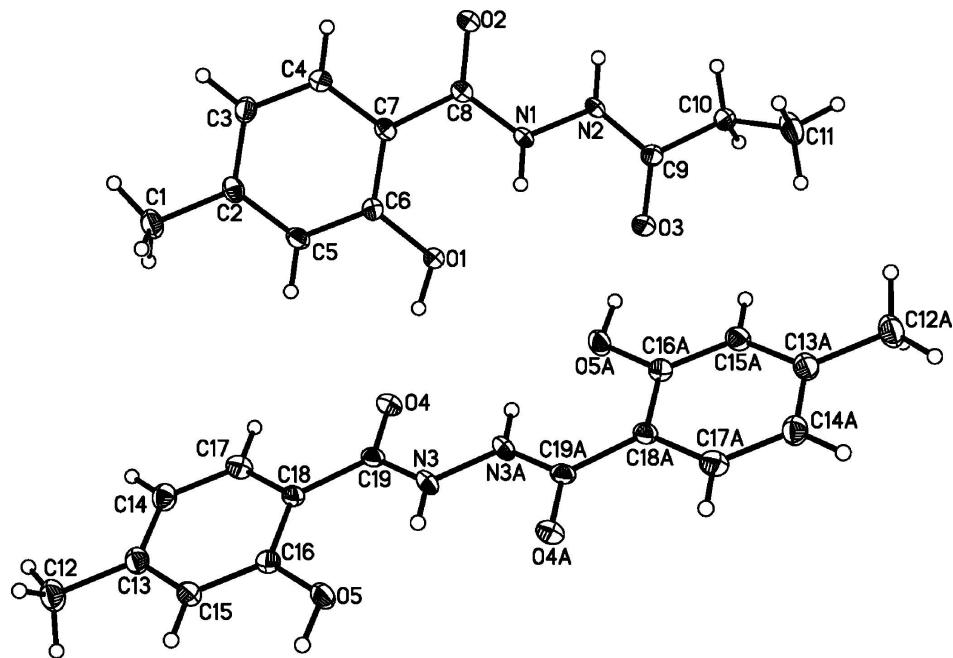
The molecular conformation is characterized by N—H···O hydrogen bonds and the crystal packing is stabilized by N—H···O and O—H···O hydrogen bonds (Fig. 2).

S2. Experimental

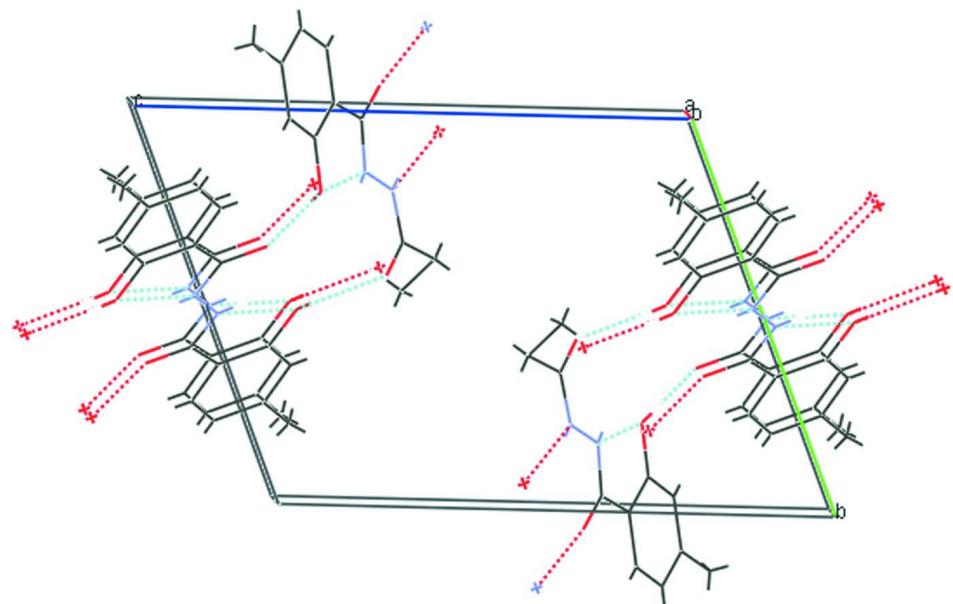
Propionic anhydride (0.26 g, 2.00 mmol) and 2-hydroxy-4-methylbenzohydrazide (0.31 g, 1.80 mmol) were stirred with an external ice-water bath in DMF (20 ml) for 6 h. The filtrate was evaporated on a rotary evaporator. After recrystallization, the title compound were obtained.

S3. Refinement

All H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms (C—H = 0.93 Å; N—H = 0.86 Å; O—H = 0.82 Å) and $U_{\text{iso}}(\text{H})$ values were set to $1.2U_{\text{eq}}(\text{C}, \text{N})$ and $1.5U_{\text{eq}}(\text{O})$.

**Figure 1**

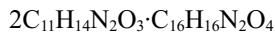
The structure of (I), showing 30% probability displacement ellipsoids and the atom-numbering scheme [symmetry code: (A) $-x + 2, -y + 1, -z$].

**Figure 2**

Packing diagram of the title compound.

2-hydroxy-4-methyl-N-propanoylbenzohydrazide– 2-hydroxy-N-(2-hydroxy-4-methylbenzoyl)-6-methylbenzohydrazide (2/1)

Crystal data



$M_r = 744.79$

Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 6.5778 (10) \text{ \AA}$

$b = 10.7618 (17) \text{ \AA}$

$c = 13.936 (2) \text{ \AA}$

$\alpha = 109.522 (3)^\circ$

$\beta = 93.608 (1)^\circ$

$\gamma = 104.448 (4)^\circ$

$V = 888.8 (2) \text{ \AA}^3$

$Z = 1$

$F(000) = 394$

$D_x = 1.391 \text{ Mg m}^{-3}$

Melting point = 488–496 K

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 6530 reflections

$\theta = 1.0\text{--}27.6^\circ$

$\mu = 0.10 \text{ mm}^{-1}$

$T = 296 \text{ K}$

Block, colourless

$0.54 \times 0.30 \times 0.25 \text{ mm}$

Data collection

Bruker APEXII

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 0 pixels mm^{-1}

ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2005)

$T_{\min} = 0.964$, $T_{\max} = 0.975$

5076 measured reflections

4125 independent reflections

1865 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.022$

$\theta_{\max} = 27.6^\circ$, $\theta_{\min} = 1.0^\circ$

$h = -8 \rightarrow 8$

$k = -14 \rightarrow 14$

$l = -18 \rightarrow 18$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.037$

$wR(F^2) = 0.110$

$S = 0.90$

4125 reflections

246 parameters

0 restraints

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

$w = 1/[\sigma^2(F_o^2) + (0.0665P)^2 + 0.5716P]$
where $P = (F_o^2 + 2F_c^2)/3$

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

$\Delta\rho_{\max} = 0.20 \text{ e \AA}^{-3}$

$\Delta\rho_{\min} = -0.22 \text{ e \AA}^{-3}$

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	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	1.5753 (4)	1.1759 (2)	0.25135 (19)	0.0371 (6)

H1A	1.6241	1.2745	0.2819	0.056*
H1B	1.5448	1.1483	0.1777	0.056*
H1C	1.6836	1.1382	0.2689	0.056*
C2	1.3767 (3)	1.1237 (2)	0.29126 (16)	0.0272 (5)
C3	1.2717 (3)	1.2119 (2)	0.35009 (16)	0.0302 (5)
H3A	1.3247	1.3062	0.3659	0.036*
C4	1.0896 (3)	1.1610 (2)	0.38542 (16)	0.0272 (5)
H4A	1.0227	1.2222	0.4251	0.033*
C5	1.2941 (3)	0.9840 (2)	0.27002 (16)	0.0271 (5)
H5A	1.3639	0.9236	0.2317	0.032*
C6	1.1100 (3)	0.9315 (2)	0.30440 (15)	0.0243 (5)
C7	1.0027 (3)	1.0199 (2)	0.36312 (15)	0.0230 (5)
C8	0.8034 (3)	0.9769 (2)	0.40407 (15)	0.0230 (5)
C9	0.4670 (3)	0.6593 (2)	0.40029 (15)	0.0272 (5)
C10	0.2783 (3)	0.6153 (2)	0.44886 (17)	0.0356 (6)
H10A	0.2494	0.6963	0.4956	0.043*
H10B	0.3117	0.5625	0.4890	0.043*
C11	0.0800 (4)	0.5283 (3)	0.3690 (2)	0.0434 (6)
H11A	-0.0362	0.5022	0.4034	0.065*
H11B	0.1071	0.4472	0.3232	0.065*
H11C	0.0442	0.5809	0.3303	0.065*
C12	2.0539 (4)	0.8165 (3)	-0.0799 (2)	0.0423 (6)
H12A	2.0282	0.8592	-0.1281	0.063*
H12B	2.1120	0.7422	-0.1122	0.063*
H12C	2.1528	0.8831	-0.0208	0.063*
C13	1.8475 (3)	0.7613 (2)	-0.04652 (17)	0.0312 (5)
C14	1.8192 (3)	0.8096 (2)	0.05601 (17)	0.0331 (5)
H14A	1.9293	0.8769	0.1054	0.040*
C15	1.6810 (3)	0.6616 (2)	-0.11824 (16)	0.0293 (5)
H15A	1.6985	0.6282	-0.1870	0.035*
C16	1.4884 (3)	0.6101 (2)	-0.08999 (16)	0.0267 (5)
C17	1.6287 (3)	0.7582 (2)	0.08499 (17)	0.0320 (5)
H17A	1.6136	0.7910	0.1541	0.038*
C18	1.4587 (3)	0.6587 (2)	0.01368 (16)	0.0257 (5)
C19	1.2600 (3)	0.6131 (2)	0.05327 (16)	0.0252 (5)
N1	0.7128 (3)	0.84114 (17)	0.37670 (13)	0.0273 (4)
H1D	0.7663	0.7838	0.3347	0.033*
N2	0.5341 (3)	0.79291 (17)	0.41560 (13)	0.0263 (4)
H2A	0.4672	0.8484	0.4492	0.032*
O4	1.2460 (2)	0.65694 (15)	0.14648 (11)	0.0333 (4)
O1	1.0312 (2)	0.79309 (14)	0.28105 (11)	0.0322 (4)
H1E	1.1084	0.7533	0.2467	0.048*
O2	0.7227 (2)	1.06102 (14)	0.45964 (11)	0.0314 (4)
O3	0.5567 (2)	0.57635 (14)	0.34881 (11)	0.0329 (4)
N3	1.0943 (3)	0.52334 (18)	-0.01587 (15)	0.0291 (5)
O5	1.3246 (2)	0.51267 (15)	-0.16289 (12)	0.0333 (4)
H3B	1.104 (4)	0.490 (2)	-0.0786 (19)	0.036 (7)*
H5B	1.368 (4)	0.481 (3)	-0.223 (2)	0.055 (8)*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0265 (12)	0.0416 (14)	0.0485 (14)	0.0077 (11)	0.0136 (10)	0.0233 (11)
C2	0.0192 (11)	0.0345 (13)	0.0299 (12)	0.0067 (10)	0.0031 (9)	0.0148 (10)
C3	0.0249 (12)	0.0268 (12)	0.0401 (13)	0.0045 (10)	0.0048 (10)	0.0158 (10)
C4	0.0249 (12)	0.0262 (12)	0.0316 (12)	0.0094 (10)	0.0067 (9)	0.0100 (9)
C5	0.0226 (12)	0.0307 (12)	0.0293 (12)	0.0108 (10)	0.0085 (9)	0.0096 (9)
C6	0.0235 (12)	0.0234 (12)	0.0256 (11)	0.0058 (9)	0.0056 (9)	0.0089 (9)
C7	0.0196 (11)	0.0280 (12)	0.0221 (11)	0.0073 (9)	0.0038 (8)	0.0095 (9)
C8	0.0208 (11)	0.0267 (12)	0.0222 (11)	0.0079 (10)	0.0042 (9)	0.0087 (9)
C9	0.0287 (12)	0.0254 (12)	0.0253 (11)	0.0071 (10)	0.0090 (9)	0.0060 (9)
C10	0.0349 (13)	0.0255 (12)	0.0442 (14)	0.0059 (10)	0.0223 (11)	0.0088 (10)
C11	0.0267 (13)	0.0530 (15)	0.0628 (17)	0.0105 (12)	0.0157 (12)	0.0356 (13)
C12	0.0302 (13)	0.0454 (14)	0.0524 (15)	0.0062 (11)	0.0121 (11)	0.0216 (12)
C13	0.0278 (12)	0.0308 (12)	0.0407 (14)	0.0104 (10)	0.0087 (10)	0.0182 (11)
C14	0.0267 (13)	0.0323 (13)	0.0373 (13)	0.0046 (10)	0.0022 (10)	0.0122 (10)
C15	0.0303 (13)	0.0303 (12)	0.0302 (12)	0.0104 (10)	0.0108 (10)	0.0123 (10)
C16	0.0259 (12)	0.0248 (11)	0.0302 (12)	0.0090 (9)	0.0061 (9)	0.0093 (9)
C17	0.0333 (13)	0.0346 (13)	0.0283 (12)	0.0126 (11)	0.0076 (10)	0.0091 (10)
C18	0.0276 (12)	0.0233 (11)	0.0313 (12)	0.0117 (9)	0.0119 (10)	0.0120 (9)
C19	0.0287 (13)	0.0233 (11)	0.0303 (13)	0.0143 (10)	0.0126 (10)	0.0117 (10)
N1	0.0245 (10)	0.0254 (10)	0.0316 (10)	0.0066 (8)	0.0167 (8)	0.0079 (8)
N2	0.0222 (10)	0.0266 (10)	0.0321 (10)	0.0089 (8)	0.0167 (8)	0.0097 (8)
O4	0.0351 (9)	0.0339 (9)	0.0309 (9)	0.0122 (7)	0.0152 (7)	0.0083 (7)
O1	0.0301 (9)	0.0232 (8)	0.0429 (9)	0.0079 (7)	0.0195 (7)	0.0087 (7)
O2	0.0284 (8)	0.0286 (8)	0.0383 (9)	0.0105 (7)	0.0155 (7)	0.0101 (7)
O3	0.0338 (9)	0.0282 (8)	0.0356 (9)	0.0094 (7)	0.0166 (7)	0.0076 (7)
N3	0.0258 (10)	0.0340 (11)	0.0290 (11)	0.0080 (9)	0.0153 (9)	0.0116 (9)
O5	0.0280 (9)	0.0366 (9)	0.0262 (9)	0.0017 (7)	0.0106 (7)	0.0044 (7)

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

C1—C2	1.505 (3)	C11—H11C	0.9600
C1—H1A	0.9600	C12—C13	1.508 (3)
C1—H1B	0.9600	C12—H12A	0.9600
C1—H1C	0.9600	C12—H12B	0.9600
C2—C5	1.385 (3)	C12—H12C	0.9600
C2—C3	1.388 (3)	C13—C15	1.382 (3)
C3—C4	1.380 (3)	C13—C14	1.390 (3)
C3—H3A	0.9300	C14—C17	1.381 (3)
C4—C7	1.399 (3)	C14—H14A	0.9300
C4—H4A	0.9300	C15—C16	1.387 (3)
C5—C6	1.388 (3)	C15—H15A	0.9300
C5—H5A	0.9300	C16—O5	1.371 (2)
C6—O1	1.365 (2)	C16—C18	1.407 (3)
C6—C7	1.400 (3)	C17—C18	1.391 (3)
C7—C8	1.492 (3)	C17—H17A	0.9300

C8—O2	1.234 (2)	C18—C19	1.485 (3)
C8—N1	1.342 (3)	C19—O4	1.243 (2)
C9—O3	1.244 (2)	C19—N3	1.330 (3)
C9—N2	1.332 (3)	N1—N2	1.382 (2)
C9—C10	1.500 (3)	N1—H1D	0.8600
C10—C11	1.526 (3)	N2—H2A	0.8600
C10—H10A	0.9700	O1—H1E	0.8200
C10—H10B	0.9700	N3—N3 ⁱ	1.376 (3)
C11—H11A	0.9600	N3—H3B	0.84 (2)
C11—H11B	0.9600	O5—H5B	0.88 (3)
C2—C1—H1A	109.5	H11A—C11—H11C	109.5
C2—C1—H1B	109.5	H11B—C11—H11C	109.5
H1A—C1—H1B	109.5	C13—C12—H12A	109.5
C2—C1—H1C	109.5	C13—C12—H12B	109.5
H1A—C1—H1C	109.5	H12A—C12—H12B	109.5
H1B—C1—H1C	109.5	C13—C12—H12C	109.5
C5—C2—C3	118.03 (19)	H12A—C12—H12C	109.5
C5—C2—C1	120.00 (19)	H12B—C12—H12C	109.5
C3—C2—C1	121.96 (19)	C15—C13—C14	118.5 (2)
C4—C3—C2	120.70 (19)	C15—C13—C12	120.2 (2)
C4—C3—H3A	119.6	C14—C13—C12	121.3 (2)
C2—C3—H3A	119.6	C17—C14—C13	120.4 (2)
C3—C4—C7	121.80 (19)	C17—C14—H14A	119.8
C3—C4—H4A	119.1	C13—C14—H14A	119.8
C7—C4—H4A	119.1	C13—C15—C16	121.5 (2)
C2—C5—C6	121.78 (19)	C13—C15—H15A	119.3
C2—C5—H5A	119.1	C16—C15—H15A	119.3
C6—C5—H5A	119.1	O5—C16—C15	120.35 (18)
O1—C6—C5	120.36 (17)	O5—C16—C18	119.40 (18)
O1—C6—C7	119.25 (17)	C15—C16—C18	120.25 (19)
C5—C6—C7	120.39 (18)	C14—C17—C18	121.9 (2)
C4—C7—C6	117.28 (18)	C14—C17—H17A	119.1
C4—C7—C8	117.01 (18)	C18—C17—H17A	119.1
C6—C7—C8	125.70 (18)	C17—C18—C16	117.49 (19)
O2—C8—N1	120.91 (18)	C17—C18—C19	117.19 (18)
O2—C8—C7	122.27 (18)	C16—C18—C19	125.33 (19)
N1—C8—C7	116.81 (17)	O4—C19—N3	120.78 (19)
O3—C9—N2	121.63 (19)	O4—C19—C18	122.20 (19)
O3—C9—C10	122.19 (18)	N3—C19—C18	117.01 (18)
N2—C9—C10	116.18 (18)	C8—N1—N2	120.37 (16)
C9—C10—C11	112.34 (18)	C8—N1—H1D	119.8
C9—C10—H10A	109.1	N2—N1—H1D	119.8
C11—C10—H10A	109.1	C9—N2—N1	119.05 (16)
C9—C10—H10B	109.1	C9—N2—H2A	120.5
C11—C10—H10B	109.1	N1—N2—H2A	120.5
H10A—C10—H10B	107.9	C6—O1—H1E	109.5
C10—C11—H11A	109.5	C19—N3—N3 ⁱ	119.9 (2)

C10—C11—H11B	109.5	C19—N3—H3B	121.4 (16)
H11A—C11—H11B	109.5	N3 ⁱ —N3—H3B	118.6 (16)
C10—C11—H11C	109.5	C16—O5—H5B	110.8 (17)

Symmetry code: (i) $-x+2, -y+1, -z$.

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1D···O1	0.86	1.93	2.620 (2)	136
O1—H1E···O4	0.82	1.87	2.682 (2)	169
N2—H2A···O2 ⁱⁱ	0.86	2.03	2.866 (2)	165
N3—H3B···O5	0.84 (2)	1.94 (3)	2.613 (3)	136 (2)
N3—H3B···O4 ⁱ	0.84 (2)	2.37 (3)	2.655 (3)	101 (2)
O5—H5B···O3 ⁱ	0.89 (3)	1.80 (3)	2.685 (2)	175 (2)

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