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5-Methyl-2-phenyl-2H-pyrazol-3-ol

Qiang Wang,^a Yi Zhang,^a Rong Wang,^a Yi-Lin Yang^{a*} and Feng Zhi^{b*}

^aDepartment of Neurosurgery, Third Affiliated Hospital of Soochow University, Changzhou 213003, People's Republic of China, and ^bLaboratory of Neuronal Injury and Protection, Third Affiliated Hospital of Soochow University, Changzhou 213003, People's Republic of China

Correspondence e-mail: czfph@yahoo.com.cn, danielzhi@yahoo.com.cn

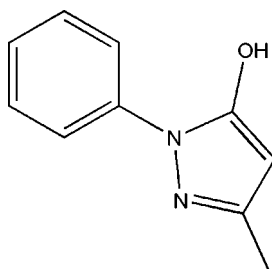
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.040; wR factor = 0.107; data-to-parameter ratio = 16.1.

The title compound, $\text{C}_{10}\text{H}_{10}\text{N}_2\text{O}$, known as Edaravone (MCI-186), was crystallized from methanol. The two independent molecules in the asymmetric unit are linked through an $\text{O}-\text{H}\cdots\text{O}$ hydrogen bond. One molecule adopts a ketone form, while the other adopts an enol form. In the crystal structure, molecules are linked through intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds, forming chains running along the b axis.

Related literature

For background to the compound, see: Watanabe *et al.* (1994); The Edaravone Acute Infarction Study Group (2003). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

 $\text{C}_{10}\text{H}_{10}\text{N}_2\text{O}$ $M_r = 174.20$

Monoclinic, $P2_1/c$
 $a = 10.336$ (2) Å
 $b = 11.154$ (2) Å
 $c = 15.863$ (3) Å
 $\beta = 95.157$ (3)°
 $V = 1821.4$ (6) Å³

$Z = 8$
 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 298$ (2) K
 $0.23 \times 0.23 \times 0.20$ mm

Data collection

Bruker SMART CCD area-detector diffractometer
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)
 $T_{\min} = 0.981$, $T_{\max} = 0.983$

10551 measured reflections
 3923 independent reflections
 2894 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.106$
 $S = 1.04$
 3923 reflections
 243 parameters
 2 restraints

H atoms treated by a mixture of independent and constrained refinement
 $\Delta\rho_{\text{max}} = 0.17$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.14$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N2}-\text{H2A}\cdots\text{N4}^i$	0.907 (9)	1.904 (10)	2.7999 (17)	169.4 (17)
$\text{O2}-\text{H2B}\cdots\text{O1}$	0.870 (9)	1.618 (10)	2.4813 (15)	170.9 (19)

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

Data collection: SMART (Bruker, 2002); cell refinement: SAINT (Bruker, 2002); 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: SHELXL97.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SG2259).

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supporting information

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5-Methyl-2-phenyl-2*H*-pyrazol-3-ol

Qiang Wang, Yi Zhang, Rong Wang, Yi-Lin Yang and Feng Zhi

S1. Comment

Edaravone (5-methyl-2-phenyl-2*H*-pyrazol-3-ol, MCI-186) is a free-radical scavenger that was approved by the Ministry of Health, Labor, and Welfare of Japan in 2001, and is now widely used for the treatment of acute cerebral infarction (Watanabe *et al.*, 1994; The Edaravone Acute Infarction Study Group, 2003). We report in this paper the crystal structure of the compound, (I).

The compound consists of two independent molecules (Fig. 1), which are linked through an intramolecular O2—H2B···O1 hydrogen bond (Table 1). One molecule adopts a ketone form, while the other adopts an enol form. In the ketone molecule, the dihedral angle between the C1—C6 benzene ring and the N1/N2/C9/C8/C7 ring is 22.9 (2)°. In the enol molecule, the dihedral angle between the C11—C16 benzene ring and the N3/N4/C19/C18/C17 ring is 34.3 (2)°. In the compound, all the bond lengths are within normal ranges (Allen *et al.*, 1987). The bond length of C17—O2 [1.325 (2) Å] is longer than that of C7—O1 [1.260 (2) Å], which is caused by the enol form.

In the crystal structure, molecules are linked through intermolecular N—H···O hydrogen bonds (Table 1), forming chains running along the *b* axis (Fig. 2).

S2. Experimental

The crystal of the compound was recrystallized by edaravone in methanol.

S3. Refinement

H2A and H2B were located in a difference Fourier map and refined isotropically, with N—H and O—H distances restrained to 0.90 (1) and 0.85 (1) Å, and with $U_{\text{iso}}(\text{H})$ set to 0.08 Å². Other H atoms were constrained to idealized geometries, with C—H = 0.93–0.96 Å, and with $U_{\text{iso}}(\text{H})$ set to 1.2 $U_{\text{eq}}(\text{C})$ and 1.5 $U_{\text{eq}}(\text{methyl C})$.

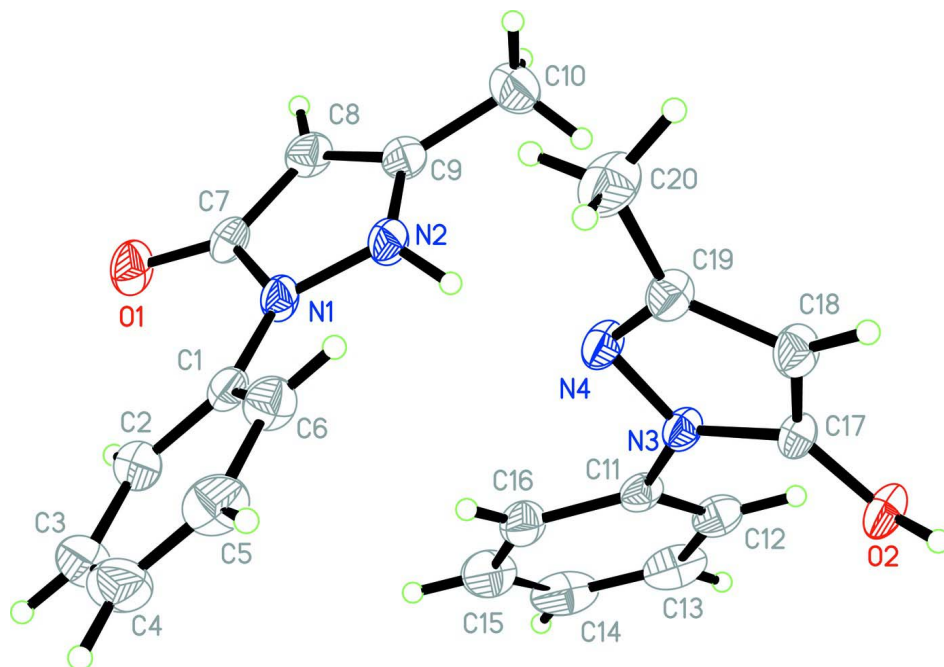


Figure 1

The structure of (I) at the 30% probability level. Intramolecular hydrogen bonds are shown as dashed lines.

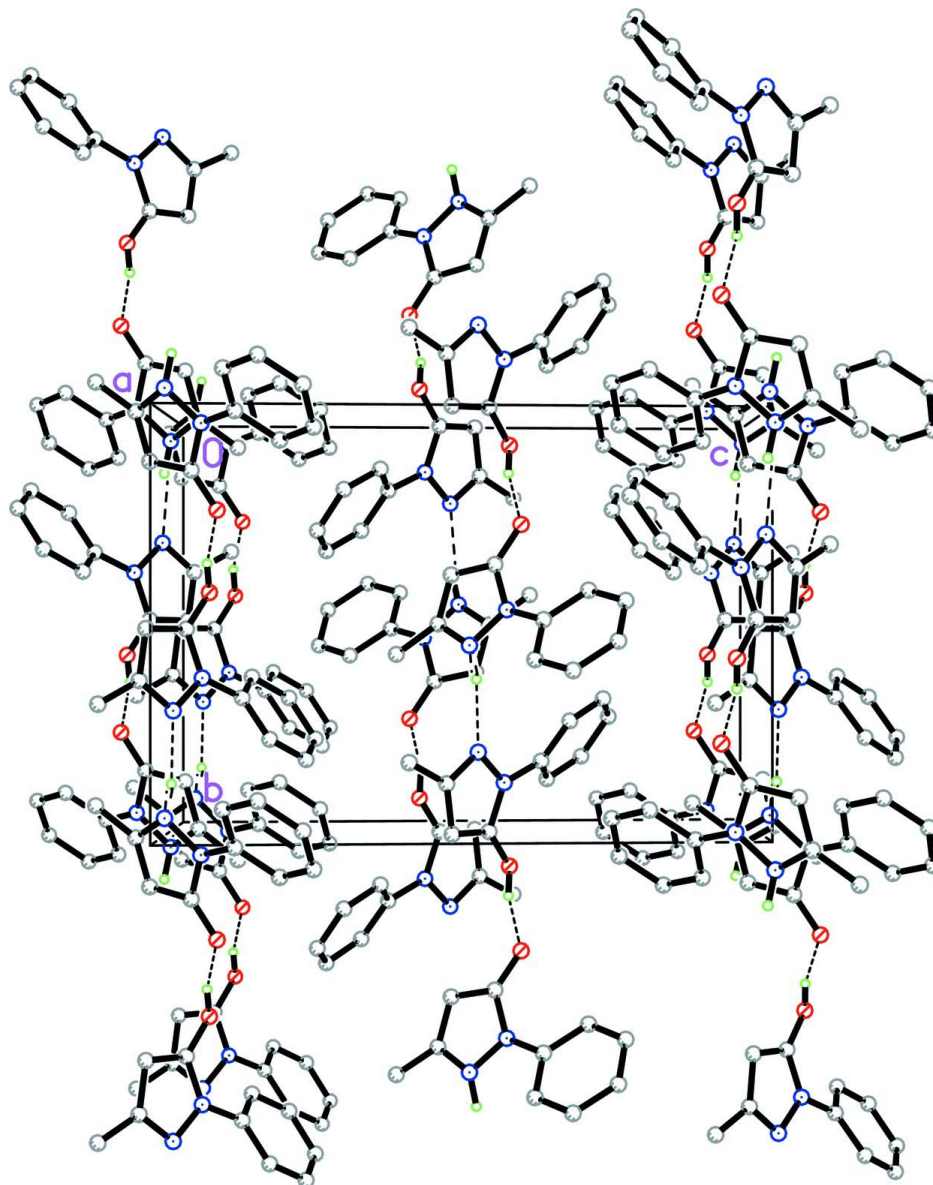


Figure 2

Molecular packing of (I), viewed along the a axis. Intermolecular hydrogen bonds are shown as dashed lines.

5-Methyl-2-phenyl-2H-pyrazol-3-ol

Crystal data

$C_{10}H_{10}N_2O$

$M_r = 174.20$

Monoclinic, $P2_1/c$

Hall symbol: $-P\ 2_1/c$

$a = 10.336\ (2)\ \text{\AA}$

$b = 11.154\ (2)\ \text{\AA}$

$c = 15.863\ (3)\ \text{\AA}$

$\beta = 95.157\ (3)^\circ$

$V = 1821.4\ (6)\ \text{\AA}^3$

$Z = 8$

$F(000) = 736$

$D_x = 1.271\ \text{Mg m}^{-3}$

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

Cell parameters from 3931 reflections

$\theta = 2.3\text{--}29.0^\circ$

$\mu = 0.09\ \text{mm}^{-1}$

$T = 298\ \text{K}$

Block, colorless

$0.23 \times 0.23 \times 0.20\ \text{mm}$

Data collection

Bruker SMART CCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scan
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.981$, $T_{\max} = 0.983$

10551 measured reflections
3923 independent reflections
2894 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$
 $\theta_{\max} = 27.0^\circ$, $\theta_{\min} = 2.2^\circ$
 $h = -13 \rightarrow 11$
 $k = -13 \rightarrow 14$
 $l = -20 \rightarrow 18$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.106$
 $S = 1.04$
3923 reflections
243 parameters
2 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
 $w = 1/[\sigma^2(F_o^2) + (0.0428P)^2 + 0.3053P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.17 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.14 \text{ e } \text{\AA}^{-3}$
Extinction correction: *SHELXL*,
 $F_c^* = kFc[1 + 0.001x\text{Fc}^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0087 (12)

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.76081 (11)	0.23408 (9)	0.09473 (8)	0.0652 (3)
O2	0.63865 (11)	0.42520 (9)	0.07652 (7)	0.0577 (3)
N1	0.80235 (11)	0.03414 (9)	0.07061 (8)	0.0469 (3)
N2	0.76341 (13)	-0.04917 (10)	0.00921 (8)	0.0516 (3)
N3	0.69321 (11)	0.62596 (9)	0.06745 (7)	0.0418 (3)
N4	0.76998 (12)	0.70070 (10)	0.02382 (8)	0.0490 (3)
C1	0.91015 (13)	0.01075 (12)	0.12978 (9)	0.0457 (3)
C2	0.92942 (16)	0.07840 (15)	0.20276 (10)	0.0603 (4)
H2	0.8722	0.1399	0.2131	0.072*
C3	1.03427 (19)	0.05375 (19)	0.26009 (12)	0.0768 (5)
H3	1.0481	0.0997	0.3090	0.092*
C4	1.1181 (2)	-0.0372 (2)	0.24600 (13)	0.0814 (6)
H4	1.1881	-0.0536	0.2853	0.098*
C5	1.09872 (17)	-0.10395 (18)	0.17403 (14)	0.0755 (5)
H5	1.1557	-0.1661	0.1647	0.091*

C6	0.99550 (15)	-0.08041 (15)	0.11475 (11)	0.0597 (4)
H6	0.9837	-0.1255	0.0653	0.072*
C7	0.73814 (14)	0.14046 (12)	0.05144 (10)	0.0486 (4)
C8	0.65398 (16)	0.11675 (14)	-0.02083 (10)	0.0566 (4)
H8	0.5970	0.1713	-0.0486	0.068*
C9	0.66976 (15)	0.00076 (14)	-0.04331 (10)	0.0536 (4)
C10	0.6047 (2)	-0.07170 (17)	-0.11335 (11)	0.0760 (5)
H10A	0.6586	-0.0744	-0.1596	0.114*
H10B	0.5227	-0.0359	-0.1321	0.114*
H10C	0.5908	-0.1517	-0.0937	0.114*
C11	0.61818 (14)	0.67394 (12)	0.12999 (8)	0.0441 (3)
C12	0.49765 (15)	0.62664 (14)	0.14196 (10)	0.0545 (4)
H12	0.4655	0.5611	0.1104	0.065*
C13	0.42556 (18)	0.67832 (19)	0.20167 (12)	0.0755 (6)
H13	0.3450	0.6462	0.2108	0.091*
C14	0.4708 (2)	0.7760 (2)	0.24752 (12)	0.0860 (7)
H14	0.4205	0.8110	0.2866	0.103*
C15	0.5904 (2)	0.82150 (17)	0.23553 (11)	0.0799 (6)
H15	0.6213	0.8876	0.2669	0.096*
C16	0.66614 (18)	0.77067 (13)	0.17737 (10)	0.0584 (4)
H16	0.7482	0.8012	0.1703	0.070*
C17	0.70530 (13)	0.51085 (11)	0.04194 (9)	0.0430 (3)
C18	0.79096 (15)	0.51055 (13)	-0.01882 (10)	0.0526 (4)
H18	0.8190	0.4445	-0.0479	0.063*
C19	0.82756 (15)	0.62954 (13)	-0.02813 (10)	0.0510 (4)
C20	0.91583 (19)	0.68181 (17)	-0.08846 (13)	0.0794 (6)
H20A	0.9060	0.7674	-0.0900	0.119*
H20B	0.8938	0.6496	-0.1440	0.119*
H20C	1.0042	0.6619	-0.0700	0.119*
H2A	0.7692 (17)	-0.1287 (9)	0.0208 (11)	0.080*
H2B	0.6823 (16)	0.3585 (12)	0.0773 (12)	0.080*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0757 (7)	0.0322 (5)	0.0864 (8)	0.0043 (5)	-0.0001 (6)	-0.0043 (5)
O2	0.0673 (7)	0.0311 (5)	0.0772 (7)	0.0007 (5)	0.0205 (6)	0.0059 (5)
N1	0.0531 (7)	0.0283 (6)	0.0577 (7)	0.0011 (5)	-0.0031 (6)	0.0011 (5)
N2	0.0640 (8)	0.0310 (6)	0.0581 (7)	0.0018 (5)	-0.0030 (6)	0.0012 (5)
N3	0.0493 (6)	0.0298 (6)	0.0476 (6)	0.0024 (5)	0.0114 (5)	0.0017 (5)
N4	0.0588 (8)	0.0324 (6)	0.0582 (7)	-0.0021 (5)	0.0185 (6)	0.0015 (5)
C1	0.0456 (8)	0.0362 (7)	0.0551 (8)	-0.0040 (6)	0.0031 (6)	0.0098 (6)
C2	0.0646 (10)	0.0555 (10)	0.0603 (10)	-0.0023 (8)	0.0035 (8)	0.0003 (8)
C3	0.0793 (13)	0.0876 (14)	0.0607 (11)	-0.0113 (11)	-0.0084 (9)	0.0024 (10)
C4	0.0653 (12)	0.1020 (16)	0.0736 (12)	-0.0004 (11)	-0.0122 (10)	0.0206 (12)
C5	0.0569 (10)	0.0769 (12)	0.0919 (14)	0.0161 (9)	0.0016 (10)	0.0187 (11)
C6	0.0567 (9)	0.0531 (9)	0.0687 (10)	0.0082 (7)	0.0024 (8)	0.0050 (8)
C7	0.0513 (8)	0.0310 (7)	0.0640 (9)	0.0023 (6)	0.0082 (7)	0.0075 (7)

C8	0.0597 (9)	0.0453 (9)	0.0635 (10)	0.0101 (7)	-0.0019 (8)	0.0127 (7)
C9	0.0599 (9)	0.0477 (9)	0.0523 (8)	0.0013 (7)	-0.0006 (7)	0.0069 (7)
C10	0.0886 (13)	0.0739 (12)	0.0622 (10)	-0.0045 (10)	-0.0120 (9)	-0.0032 (9)
C11	0.0544 (8)	0.0339 (7)	0.0447 (7)	0.0122 (6)	0.0093 (6)	0.0067 (6)
C12	0.0524 (9)	0.0531 (9)	0.0586 (9)	0.0130 (7)	0.0090 (7)	0.0135 (7)
C13	0.0686 (11)	0.0872 (14)	0.0749 (12)	0.0326 (10)	0.0304 (9)	0.0286 (11)
C14	0.1171 (18)	0.0853 (15)	0.0612 (11)	0.0533 (13)	0.0383 (12)	0.0143 (11)
C15	0.1298 (18)	0.0540 (11)	0.0579 (10)	0.0258 (11)	0.0198 (11)	-0.0063 (8)
C16	0.0812 (11)	0.0414 (8)	0.0537 (9)	0.0066 (8)	0.0121 (8)	-0.0019 (7)
C17	0.0495 (8)	0.0285 (6)	0.0511 (7)	0.0027 (6)	0.0048 (6)	0.0015 (6)
C18	0.0622 (9)	0.0379 (8)	0.0597 (9)	0.0065 (7)	0.0166 (7)	-0.0068 (7)
C19	0.0538 (8)	0.0432 (8)	0.0580 (9)	0.0022 (6)	0.0163 (7)	-0.0002 (7)
C20	0.0868 (13)	0.0683 (12)	0.0901 (13)	-0.0032 (10)	0.0467 (11)	0.0008 (10)

Geometric parameters (Å, °)

O1—C7	1.260 (2)	C8—C9	1.356 (2)
O2—C17	1.325 (2)	C8—H8	0.9300
O2—H2B	0.870 (9)	C9—C10	1.485 (2)
N1—C7	1.379 (2)	C10—H10A	0.9600
N1—N2	1.380 (2)	C10—H10B	0.9600
N1—C1	1.415 (2)	C10—H10C	0.9600
N2—C9	1.340 (2)	C11—C16	1.381 (2)
N2—H2A	0.907 (9)	C11—C12	1.382 (2)
N3—C17	1.355 (2)	C12—C13	1.383 (2)
N3—N4	1.379 (2)	C12—H12	0.9300
N3—C11	1.418 (2)	C13—C14	1.369 (3)
N4—C19	1.323 (2)	C13—H13	0.9300
C1—C6	1.381 (2)	C14—C15	1.365 (3)
C1—C2	1.381 (2)	C14—H14	0.9300
C2—C3	1.378 (2)	C15—C16	1.384 (2)
C2—H2	0.9300	C15—H15	0.9300
C3—C4	1.366 (3)	C16—H16	0.9300
C3—H3	0.9300	C17—C18	1.366 (2)
C4—C5	1.362 (3)	C18—C19	1.392 (2)
C4—H4	0.9300	C18—H18	0.9300
C5—C6	1.383 (2)	C19—C20	1.498 (2)
C5—H5	0.9300	C20—H20A	0.9600
C6—H6	0.9300	C20—H20B	0.9600
C7—C8	1.400 (2)	C20—H20C	0.9600
C17—O2—H2B	109.5 (13)	C9—C10—H10B	109.5
C7—N1—N2	108.62 (11)	H10A—C10—H10B	109.5
C7—N1—C1	129.74 (12)	C9—C10—H10C	109.5
N2—N1—C1	120.42 (11)	H10A—C10—H10C	109.5
C9—N2—N1	107.91 (11)	H10B—C10—H10C	109.5
C9—N2—H2A	124.3 (11)	C16—C11—C12	120.50 (14)
N1—N2—H2A	120.4 (12)	C16—C11—N3	118.91 (13)

C17—N3—N4	110.49 (11)	C12—C11—N3	120.57 (13)
C17—N3—C11	129.60 (11)	C11—C12—C13	118.88 (17)
N4—N3—C11	119.90 (10)	C11—C12—H12	120.6
C19—N4—N3	105.14 (11)	C13—C12—H12	120.6
C6—C1—C2	120.05 (14)	C14—C13—C12	121.09 (19)
C6—C1—N1	119.90 (13)	C14—C13—H13	119.5
C2—C1—N1	120.05 (13)	C12—C13—H13	119.5
C3—C2—C1	119.32 (17)	C15—C14—C13	119.52 (17)
C3—C2—H2	120.3	C15—C14—H14	120.2
C1—C2—H2	120.3	C13—C14—H14	120.2
C4—C3—C2	120.86 (18)	C14—C15—C16	120.93 (19)
C4—C3—H3	119.6	C14—C15—H15	119.5
C2—C3—H3	119.6	C16—C15—H15	119.5
C5—C4—C3	119.69 (17)	C11—C16—C15	119.06 (18)
C5—C4—H4	120.2	C11—C16—H16	120.5
C3—C4—H4	120.2	C15—C16—H16	120.5
C4—C5—C6	120.84 (18)	O2—C17—N3	119.60 (12)
C4—C5—H5	119.6	O2—C17—C18	133.19 (13)
C6—C5—H5	119.6	N3—C17—C18	107.21 (12)
C1—C6—C5	119.23 (17)	C17—C18—C19	105.80 (12)
C1—C6—H6	120.4	C17—C18—H18	127.1
C5—C6—H6	120.4	C19—C18—H18	127.1
O1—C7—N1	122.01 (13)	N4—C19—C18	111.36 (13)
O1—C7—C8	132.37 (13)	N4—C19—C20	119.87 (13)
N1—C7—C8	105.62 (12)	C18—C19—C20	128.74 (14)
C9—C8—C7	108.35 (13)	C19—C20—H20A	109.5
C9—C8—H8	125.8	C19—C20—H20B	109.5
C7—C8—H8	125.8	H20A—C20—H20B	109.5
N2—C9—C8	109.29 (14)	C19—C20—H20C	109.5
N2—C9—C10	119.47 (14)	H20A—C20—H20C	109.5
C8—C9—C10	131.23 (15)	H20B—C20—H20C	109.5
C9—C10—H10A	109.5		

Hydrogen-bond geometry (Å, °)

<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>
N2—H2 <i>A</i> ...N4 ⁱ	0.91 (1)	1.90 (1)	2.7999 (17)	169 (2)
O2—H2 <i>B</i> ...O1	0.87 (1)	1.62 (1)	2.4813 (15)	171 (2)

Symmetry code: (i) *x*, *y*-1, *z*.