

Ethyl 4-(3-hydroxyphenyl)-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline-3-carboxylate

P. Mookiah,^a K. Rajesh,^b T. Narasimhamurthy,^c
V. Vijayakumar^d and N. Srinivasan^{a*}

^aDepartment of Physics, Thiagarajar College, Madurai 625 009, India, ^bOrganic Chemistry Division, School of Science, VIT University, Vellore 632 014, India, ^cMaterials Research Centre, Indian Institute of Science, Bangalore 560 012, India, and ^dOrganic Chemistry Division, School of Science, VIT University, Vellore 632 014, India

Correspondence e-mail: vasan692000@yahoo.co.in

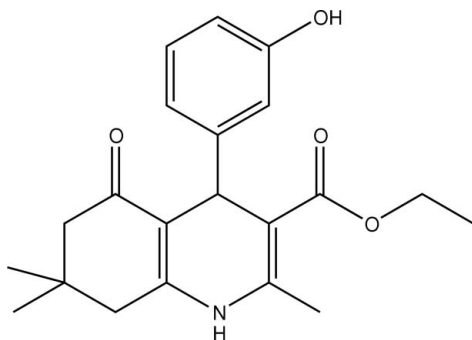
Received 17 September 2009; accepted 30 September 2009

Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; R factor = 0.040; wR factor = 0.115; data-to-parameter ratio = 13.4.

In the molecular structure of the title compound, $\text{C}_{21}\text{H}_{25}\text{NO}_4$, the dihydropyridine ring adopts a flattened boat conformation while the cyclohexenone ring is in an envelope conformation. In the crystal structure, molecules are linked into a two-dimensional network parallel to $(10\bar{1})$ by $\text{N}-\text{H}\cdots\text{O}$ and $\text{O}-\text{H}\cdots\text{O}$ hydrogen bonds. The network is generated by $R_4^4(30)$ and $R_4^4(34)$ graph-set motifs.

Related literature

For general background to oxoquinoline derivatives, see: Baba (1997); Baba *et al.* (1997,1998); Koga *et al.* (1980); Qi *et al.* (2007). For a related structure, see: Czaun *et al.* (2002); For graph-set motifs, see: Etter *et al.* (1990).



Experimental

Crystal data

$\text{C}_{21}\text{H}_{25}\text{NO}_4$

$M_r = 355.42$

Monoclinic, $P2_1/n$
 $a = 10.8721$ (4) Å
 $b = 16.1255$ (7) Å
 $c = 11.0856$ (4) Å
 $\beta = 100.682$ (2)°
 $V = 1909.83$ (13) Å³

$Z = 4$
Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 296$ K
 $0.26 \times 0.15 \times 0.12$ mm

Data collection

Bruker Kappa APEXII area-detector diffractometer
Absorption correction: multi-scan (SADABS; Sheldrick, 2004)
 $T_{\min} = 0.93$, $T_{\max} = 0.95$

14667 measured reflections
3163 independent reflections
2137 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.041$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.115$
 $S = 1.02$
3163 reflections

236 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.15$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.15$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|--|-------|-------------|-------------|---------------|
| $\text{O8C}-\text{H8C}\cdots\text{O9B}^{\text{i}}$ | 0.82 | 2.05 | 2.835 (2) | 162 |
| $\text{N1}-\text{H1}\cdots\text{O6A}^{\text{ii}}$ | 0.86 | 2.16 | 2.970 (2) | 157 |

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

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

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: CI2916).

References

- Baba, M. (1997). *Antivir. Res.* **33**, 141–152.
Baba, M., Okamoto, M., Kawamura, M., Makino, M., Higashida, T., Takashi, T., Kimura, Y., Ikeuchi, T., Tetsuka, T. & Okamoto, T. (1998). *Mol. Pharm.* **53**, 1097–1103.
Baba, M., Okamoto, M., Makino, M., Kimura, Y., Ikeuchi, T., Sakaguchi, T. & Okamoto, T. (1997). *Antimicrob. Agents Chemoth.* **41**, 1250–1255.
Bruker (2004). APEX2 and SAINT-Plus. Bruker AXS Inc., Madison, Wisconsin, USA.
Czaun, M., Ganszky, I., Speier, G. & Parkanyi, L. (2002). *Z. Kristallogr. New Cryst. Struct.* **217**, 379–380.
Etter, M. C., MacDonald, J. C. & Bernstein, J. (1990). *Acta Cryst.* **B46**, 256–262.
Koga, H., Itoh, A., Murayama, S., Suzue, S. & Irikura, T. (1980). *J. Med. Chem.* **23**, 1358–1363.
Qi, R., Fetzner, S. & Oakley, A. J. (2007). *Acta Cryst.* **F63**, 378–381.
Sheldrick, G. M. (2004). SADABS. University of Göttingen, Germany.
Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
Spek, A. L. (2009). *Acta Cryst.* **D65**, 148–155.

supplementary materials

Acta Cryst. (2009). E65, o2664 [doi:10.1107/S1600536809039877]

Ethyl 4-(3-hydroxyphenyl)-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline-3-carboxylate

P. Mookiah, K. Rajesh, T. Narasimhamurthy, V. Vijayakumar and N. Srinivasan

Comment

Some oxoquinoline derivatives *viz.* 8-difluoromethoxy-1-ethyl-6-fluoro-1,4-dihydro-7-[4-(2-methoxyphenyl)-1-piperazinyl]-4-oxoquinoline-3-carboxylic acid (K-12), 7-(3,4-dehydro-4-phenyl-1-piperidiny)-1,4-dihydro-6-fluoro-1-methyl-8-trifluoromethyl-4-oxoquinoline-3-carboxylic acid (K-37), 8-difluoromethoxy-1,4-dihydro-6-fluoro-7-(3,4-dehydro-4-phenyl-1-piperidiny)-1-[4,(1,2,4-triazol-1-yl)methylphenyl]-4-oxoquinoline-3-carboxylic acid (K-38) act as potent and selective inhibitor of human immunodeficiency virus type I (HIV-1) transcription (Baba, 1997; Baba *et al.*, 1997,1998). Structure-activity relationships of antibacterial oxoquinolone-3-carboxylic acids have been studied (Koga *et al.*, 1980). In view of the significant biological activity, precise single crystal structure determinations of these derivatives are expected to provide insights in their design and function. The crystal structure of 1*H*-2-phenyl-3-hydroxy-4-oxoquinoline-dimethylsulfoxide has already been reported (Czaun *et al.*, 2002). The expression, purification and crystallization of 1*H*-3-hydroxy-4-oxoquinoline 2,4-dioxygenase are reported elsewhere (Qi *et al.*, 2007).

The dihydropyridine ring of the title molecule (Fig.1) adopts a flattened boat conformation. The cyclohexenone ring is in an envelope conformation with atom C4 at the flap. The 4-methoxyphenyl ring and the planar part of the dihydropyridine ring (C2/C7/C9/C10) are nearly perpendicular to each other, with a dihedral angle of 89.37 (6)°.

In the crystal structure, molecules are linked into a two-dimensional network (Fig.2) parallel to the (10 $\bar{1}$) by N—H \cdots O and O—H \cdots O hydrogen bonds (Table 1). The two-dimensional layer, resembling a corrugated sheet, contains $R_4^4(30)$ and $R_4^4(34)$ graph-set motifs (Etter *et al.*, 1990) as its fundamental repeating units. It is observed that these rings are assembled through centrosymmetrically related pairs of molecules with no direct hydrogen bonding between them.

Experimental

A 50 ml round-bottomed flask was charged with 3-hydroxybenzaldehyde (1.221 g, 10 mmol), 5,5-dimethyl-1,3-cyclohexanedione (1.402 g, 10 mmol), ethyl acetoacetate (1.265 ml, 10 mmol) and ammonium acetate (0.771 g, 10 mmol) followed by ethanol (10 ml). The mixture was stirred at 343 K for 1.5 h and left aside for a day. The solid separated out was filtered and washed with ethanol-diethyl ether mixture (1:4). It was recrystallized from 100% chloroform. Light yellow prismatic crystals of the title compound were obtained by slow evaporation of a methanolic solution. Pale yellow crystals with slab morphology were obtained by slow evaporation of a methanol-chloroform solution.

Refinement

H atoms were positioned geometrically [O-H = 0.82 Å, N-H = 0.86 Å and C-H = 0.93–0.98 Å] and refined using a riding model with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$ and $1.2U_{\text{eq}}(\text{O and C}_{\text{methyl}})$.

Figures

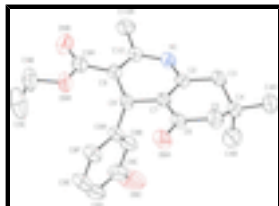


Fig. 1. The molecular structure of the title compound, with atom labels and 50% probability displacement ellipsoids for non-H atoms. H atoms have been omitted for clarity.

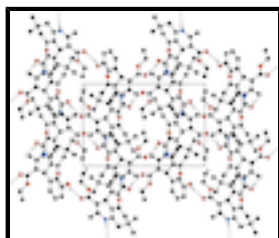


Fig. 2. A view of the molecular aggregation down the *a* axis. Hydrogen bonds are shown as dashed lines. C-bound H atoms have been omitted for clarity.

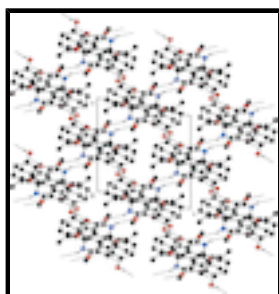


Fig. 3. A view of the molecular aggregation down the *b* axis. Hydrogen bonds are shown as dashed lines. C-bound H atoms have been omitted for clarity.

Ethyl 4-(3-hydroxyphenyl)-2,7,7-trimethyl-5-oxo-1,4,5,6,7,8-hexahydroquinoline- 3-carboxylate

Crystal data

$C_{21}H_{25}NO_4$

$M_r = 355.42$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 10.8721$ (4) Å

$b = 16.1255$ (7) Å

$c = 11.0856$ (4) Å

$\beta = 100.682$ (2)°

$V = 1909.83$ (13) Å³

$Z = 4$

$F_{000} = 760$

$D_x = 1.236$ Mg m⁻³

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

Cell parameters from 5123 reflections

$\theta = 2.0$ – 30.0 °

$\mu = 0.09$ mm⁻¹

$T = 296$ K

Prism, yellow

$0.26 \times 0.15 \times 0.12$ mm

Data collection

Bruker Kappa APEXII area-detector diffractometer

Radiation source: fine-focus sealed tube

Monochromator: graphite

$T = 296$ K

3163 independent reflections

2137 reflections with $I > 2\sigma(I)$

$R_{int} = 0.041$

$\theta_{max} = 24.5$ °

| | |
|--|-----------------------------|
| ω and φ scans | $\theta_{\min} = 2.3^\circ$ |
| Absorption correction: multi-scan (SADABS; Sheldrick, 2004) | $h = -12 \rightarrow 12$ |
| $T_{\min} = 0.93$, $T_{\max} = 0.95$ | $k = -18 \rightarrow 17$ |
| 14667 measured reflections | $l = -11 \rightarrow 12$ |

Refinement

| | |
|--|--|
| Refinement on F^2 | Secondary atom site location: difference Fourier map |
| Least-squares matrix: full | Hydrogen site location: inferred from neighbouring sites |
| $R[F^2 > 2\sigma(F^2)] = 0.040$ | H-atom parameters constrained |
| $wR(F^2) = 0.115$ | $w = 1/[\sigma^2(F_o^2) + (0.048P)^2 + 0.4479P]$ |
| $S = 1.02$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| 3163 reflections | $(\Delta/\sigma)_{\max} = 0.001$ |
| 236 parameters | $\Delta\rho_{\max} = 0.15 \text{ e } \text{\AA}^{-3}$ |
| Primary atom site location: structure-invariant direct methods | $\Delta\rho_{\min} = -0.15 \text{ e } \text{\AA}^{-3}$ |
| | Extinction correction: none |

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 | $U_{\text{iso}}^*/U_{\text{eq}}$ |
|-----|--------------|--------------|--------------|----------------------------------|
| O9A | 1.01275 (14) | 0.04777 (9) | 0.71484 (12) | 0.0559 (4) |
| O6A | 1.21908 (13) | 0.29658 (10) | 0.59297 (11) | 0.0577 (4) |
| O9B | 0.84079 (15) | -0.01070 (9) | 0.60571 (13) | 0.0618 (4) |
| O8C | 0.72595 (19) | 0.37790 (12) | 0.71929 (17) | 0.0969 (7) |
| H8C | 0.7098 | 0.4009 | 0.7804 | 0.145* |
| N1 | 0.85621 (15) | 0.18610 (10) | 0.34959 (13) | 0.0444 (4) |
| H1 | 0.7990 | 0.1933 | 0.2857 | 0.053* |
| C7 | 1.03735 (16) | 0.23880 (11) | 0.47694 (14) | 0.0346 (4) |
| C2 | 0.95697 (17) | 0.23874 (11) | 0.36798 (15) | 0.0365 (4) |
| C8 | 1.01637 (17) | 0.18488 (12) | 0.58296 (15) | 0.0376 (5) |
| H8 | 1.0962 | 0.1583 | 0.6176 | 0.045* |
| C9 | 0.92188 (17) | 0.11633 (11) | 0.53752 (15) | 0.0368 (4) |
| C6 | 1.14473 (17) | 0.29327 (12) | 0.49410 (15) | 0.0394 (5) |

supplementary materials

| | | | | |
|------|--------------|---------------|--------------|-------------|
| C10 | 0.84203 (17) | 0.12208 (12) | 0.42851 (15) | 0.0384 (5) |
| C9A | 0.91749 (19) | 0.04511 (13) | 0.61867 (17) | 0.0433 (5) |
| C4 | 1.04698 (18) | 0.37054 (12) | 0.29899 (16) | 0.0435 (5) |
| C8B | 0.8704 (2) | 0.28573 (13) | 0.65967 (17) | 0.0509 (6) |
| H8B | 0.8260 | 0.2893 | 0.5796 | 0.061* |
| C3 | 0.97318 (19) | 0.29188 (13) | 0.26167 (15) | 0.0471 (5) |
| H3A | 0.8913 | 0.3068 | 0.2159 | 0.057* |
| H3B | 1.0157 | 0.2600 | 0.2075 | 0.057* |
| C5 | 1.16617 (19) | 0.34564 (14) | 0.38772 (17) | 0.0537 (6) |
| H5A | 1.2199 | 0.3154 | 0.3424 | 0.064* |
| H5B | 1.2103 | 0.3956 | 0.4196 | 0.064* |
| C8A | 0.97467 (19) | 0.23549 (12) | 0.68484 (15) | 0.0421 (5) |
| C10A | 0.7372 (2) | 0.06394 (14) | 0.37939 (18) | 0.0536 (6) |
| H10A | 0.7707 | 0.0101 | 0.3675 | 0.080* |
| H10B | 0.6924 | 0.0845 | 0.3024 | 0.080* |
| H10C | 0.6813 | 0.0600 | 0.4368 | 0.080* |
| C8F | 1.0398 (2) | 0.23138 (15) | 0.80510 (17) | 0.0608 (6) |
| H8F | 1.1111 | 0.1986 | 0.8241 | 0.073* |
| C8C | 0.8307 (2) | 0.33082 (14) | 0.7515 (2) | 0.0608 (6) |
| C9B | 1.0157 (2) | -0.01484 (16) | 0.8081 (2) | 0.0682 (7) |
| H91B | 1.0250 | -0.0694 | 0.7740 | 0.082* |
| H92B | 0.9384 | -0.0139 | 0.8400 | 0.082* |
| C4A | 1.0813 (2) | 0.41285 (15) | 0.18658 (18) | 0.0673 (7) |
| H41A | 1.1279 | 0.4625 | 0.2115 | 0.101* |
| H42A | 1.0063 | 0.4268 | 0.1298 | 0.101* |
| H43A | 1.1313 | 0.3759 | 0.1477 | 0.101* |
| C8E | 0.9988 (3) | 0.27565 (18) | 0.8960 (2) | 0.0777 (8) |
| H8E | 1.0422 | 0.2714 | 0.9763 | 0.093* |
| C8D | 0.8956 (3) | 0.32578 (17) | 0.8710 (2) | 0.0717 (8) |
| H8D | 0.8697 | 0.3559 | 0.9333 | 0.086* |
| C4B | 0.9704 (3) | 0.43009 (15) | 0.3613 (2) | 0.0796 (8) |
| H41B | 1.0181 | 0.4796 | 0.3844 | 0.119* |
| H42B | 0.9502 | 0.4043 | 0.4332 | 0.119* |
| H43B | 0.8947 | 0.4440 | 0.3056 | 0.119* |
| C9C | 1.1234 (3) | 0.0034 (2) | 0.9074 (2) | 0.0993 (11) |
| H91C | 1.1276 | -0.0374 | 0.9710 | 0.149* |
| H92C | 1.1130 | 0.0574 | 0.9407 | 0.149* |
| H93C | 1.1993 | 0.0022 | 0.8749 | 0.149* |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|-------------|-------------|-------------|-------------|-------------|--------------|
| O9A | 0.0679 (10) | 0.0550 (9) | 0.0430 (7) | 0.0030 (7) | 0.0057 (7) | 0.0178 (7) |
| O6A | 0.0501 (9) | 0.0749 (11) | 0.0390 (7) | -0.0106 (8) | -0.0157 (7) | 0.0038 (7) |
| O9B | 0.0681 (11) | 0.0521 (10) | 0.0667 (10) | -0.0076 (8) | 0.0164 (8) | 0.0124 (8) |
| O8C | 0.1049 (15) | 0.1028 (16) | 0.0858 (13) | 0.0344 (13) | 0.0246 (11) | -0.0337 (11) |
| N1 | 0.0460 (10) | 0.0493 (10) | 0.0313 (8) | -0.0099 (8) | -0.0099 (7) | 0.0028 (8) |
| C7 | 0.0365 (10) | 0.0380 (11) | 0.0272 (9) | 0.0017 (8) | 0.0006 (7) | -0.0014 (8) |

| | | | | | | |
|------|-------------|-------------|-------------|--------------|--------------|--------------|
| C2 | 0.0402 (11) | 0.0380 (11) | 0.0286 (9) | -0.0024 (9) | -0.0005 (8) | -0.0028 (8) |
| C8 | 0.0390 (11) | 0.0427 (11) | 0.0282 (9) | 0.0027 (9) | -0.0011 (8) | 0.0036 (8) |
| C9 | 0.0438 (11) | 0.0356 (11) | 0.0325 (9) | 0.0021 (9) | 0.0107 (8) | -0.0012 (8) |
| C6 | 0.0375 (11) | 0.0456 (12) | 0.0312 (9) | 0.0008 (9) | -0.0034 (8) | -0.0018 (9) |
| C10 | 0.0438 (11) | 0.0374 (11) | 0.0338 (9) | -0.0024 (9) | 0.0068 (8) | -0.0046 (9) |
| C9A | 0.0479 (12) | 0.0432 (12) | 0.0418 (11) | 0.0067 (10) | 0.0160 (10) | 0.0002 (9) |
| C4 | 0.0519 (12) | 0.0440 (12) | 0.0312 (9) | -0.0051 (10) | -0.0009 (9) | 0.0020 (9) |
| C8B | 0.0619 (14) | 0.0548 (14) | 0.0351 (10) | 0.0008 (11) | 0.0063 (10) | -0.0091 (10) |
| C3 | 0.0542 (13) | 0.0540 (13) | 0.0282 (9) | -0.0119 (10) | -0.0052 (9) | 0.0032 (9) |
| C5 | 0.0508 (13) | 0.0627 (14) | 0.0429 (11) | -0.0151 (11) | -0.0037 (10) | 0.0060 (10) |
| C8A | 0.0538 (13) | 0.0429 (12) | 0.0285 (9) | -0.0071 (10) | 0.0046 (8) | -0.0019 (8) |
| C10A | 0.0579 (14) | 0.0541 (14) | 0.0472 (11) | -0.0153 (11) | 0.0058 (10) | -0.0075 (10) |
| C8F | 0.0772 (16) | 0.0680 (15) | 0.0326 (11) | -0.0053 (13) | -0.0016 (10) | -0.0034 (11) |
| C8C | 0.0738 (16) | 0.0540 (15) | 0.0586 (14) | -0.0031 (13) | 0.0229 (12) | -0.0155 (12) |
| C9B | 0.0835 (18) | 0.0674 (16) | 0.0571 (13) | 0.0206 (13) | 0.0220 (13) | 0.0296 (12) |
| C4A | 0.0850 (18) | 0.0689 (16) | 0.0438 (12) | -0.0258 (14) | 0.0008 (11) | 0.0110 (11) |
| C8E | 0.109 (2) | 0.089 (2) | 0.0303 (11) | -0.0111 (18) | 0.0015 (13) | -0.0135 (12) |
| C8D | 0.101 (2) | 0.0724 (18) | 0.0468 (13) | -0.0194 (16) | 0.0261 (14) | -0.0260 (13) |
| C4B | 0.117 (2) | 0.0556 (16) | 0.0675 (15) | 0.0266 (15) | 0.0204 (15) | 0.0067 (13) |
| C9C | 0.0805 (19) | 0.148 (3) | 0.0679 (16) | 0.0255 (19) | 0.0092 (15) | 0.0560 (19) |

Geometric parameters (Å, °)

| | | | |
|----------|-----------|-----------|-----------|
| O9A—C9A | 1.342 (2) | C3—H3A | 0.97 |
| O9A—C9B | 1.442 (2) | C3—H3B | 0.97 |
| O6A—C6 | 1.237 (2) | C5—H5A | 0.97 |
| O9B—C9A | 1.217 (2) | C5—H5B | 0.97 |
| O8C—C8C | 1.361 (3) | C8A—C8F | 1.390 (3) |
| O8C—H8C | 0.82 | C10A—H10A | 0.96 |
| N1—C2 | 1.371 (2) | C10A—H10B | 0.96 |
| N1—C10 | 1.380 (2) | C10A—H10C | 0.96 |
| N1—H1 | 0.86 | C8F—C8E | 1.374 (3) |
| C7—C2 | 1.353 (2) | C8F—H8F | 0.93 |
| C7—C6 | 1.445 (3) | C8C—C8D | 1.383 (3) |
| C7—C8 | 1.513 (2) | C9B—C9C | 1.480 (3) |
| C2—C3 | 1.494 (3) | C9B—H91B | 0.97 |
| C8—C9 | 1.529 (3) | C9B—H92B | 0.97 |
| C8—C8A | 1.529 (3) | C4A—H41A | 0.96 |
| C8—H8 | 0.98 | C4A—H42A | 0.96 |
| C9—C10 | 1.354 (2) | C4A—H43A | 0.96 |
| C9—C9A | 1.465 (3) | C8E—C8D | 1.369 (4) |
| C6—C5 | 1.503 (3) | C8E—H8E | 0.93 |
| C10—C10A | 1.498 (3) | C8D—H8D | 0.93 |
| C4—C3 | 1.517 (3) | C4B—H41B | 0.96 |
| C4—C4B | 1.518 (3) | C4B—H42B | 0.96 |
| C4—C4A | 1.526 (3) | C4B—H43B | 0.96 |
| C4—C5 | 1.528 (3) | C9C—H91C | 0.96 |
| C8B—C8A | 1.379 (3) | C9C—H92C | 0.96 |
| C8B—C8C | 1.383 (3) | C9C—H93C | 0.96 |

supplementary materials

| | | | |
|-------------|-------------|----------------|-------------|
| C8B—H8B | 0.93 | | |
| C9A—O9A—C9B | 117.31 (17) | C4—C5—H5B | 108.6 |
| C8C—O8C—H8C | 109.5 | H5A—C5—H5B | 107.6 |
| C2—N1—C10 | 123.30 (14) | C8B—C8A—C8F | 118.38 (19) |
| C2—N1—H1 | 118.4 | C8B—C8A—C8 | 120.62 (15) |
| C10—N1—H1 | 118.4 | C8F—C8A—C8 | 120.99 (19) |
| C2—C7—C6 | 119.35 (16) | C10—C10A—H10A | 109.5 |
| C2—C7—C8 | 121.84 (17) | C10—C10A—H10B | 109.5 |
| C6—C7—C8 | 118.79 (14) | H10A—C10A—H10B | 109.5 |
| C7—C2—N1 | 119.95 (17) | C10—C10A—H10C | 109.5 |
| C7—C2—C3 | 123.56 (17) | H10A—C10A—H10C | 109.5 |
| N1—C2—C3 | 116.47 (14) | H10B—C10A—H10C | 109.5 |
| C7—C8—C9 | 110.38 (14) | C8E—C8F—C8A | 120.1 (2) |
| C7—C8—C8A | 112.04 (15) | C8E—C8F—H8F | 120.0 |
| C9—C8—C8A | 110.86 (15) | C8A—C8F—H8F | 120.0 |
| C7—C8—H8 | 107.8 | O8C—C8C—C8B | 117.4 (2) |
| C9—C8—H8 | 107.8 | O8C—C8C—C8D | 122.5 (2) |
| C8A—C8—H8 | 107.8 | C8B—C8C—C8D | 120.1 (2) |
| C10—C9—C9A | 120.87 (17) | O9A—C9B—C9C | 107.7 (2) |
| C10—C9—C8 | 121.66 (16) | O9A—C9B—H91B | 110.2 |
| C9A—C9—C8 | 117.43 (15) | C9C—C9B—H91B | 110.2 |
| O6A—C6—C7 | 121.49 (17) | O9A—C9B—H92B | 110.2 |
| O6A—C6—C5 | 120.01 (17) | C9C—C9B—H92B | 110.2 |
| C7—C6—C5 | 118.50 (14) | H91B—C9B—H92B | 108.5 |
| C9—C10—N1 | 119.23 (16) | C4—C4A—H41A | 109.5 |
| C9—C10—C10A | 126.89 (18) | C4—C4A—H42A | 109.5 |
| N1—C10—C10A | 113.86 (15) | H41A—C4A—H42A | 109.5 |
| O9B—C9A—O9A | 121.89 (18) | C4—C4A—H43A | 109.5 |
| O9B—C9A—C9 | 127.35 (18) | H41A—C4A—H43A | 109.5 |
| O9A—C9A—C9 | 110.76 (17) | H42A—C4A—H43A | 109.5 |
| C3—C4—C4B | 110.26 (19) | C8D—C8E—C8F | 121.6 (2) |
| C3—C4—C4A | 110.37 (15) | C8D—C8E—H8E | 119.2 |
| C4B—C4—C4A | 109.04 (18) | C8F—C8E—H8E | 119.2 |
| C3—C4—C5 | 107.36 (16) | C8E—C8D—C8C | 118.8 (2) |
| C4B—C4—C5 | 110.14 (17) | C8E—C8D—H8D | 120.6 |
| C4A—C4—C5 | 109.66 (17) | C8C—C8D—H8D | 120.6 |
| C8A—C8B—C8C | 121.09 (19) | C4—C4B—H41B | 109.5 |
| C8A—C8B—H8B | 119.5 | C4—C4B—H42B | 109.5 |
| C8C—C8B—H8B | 119.5 | H41B—C4B—H42B | 109.5 |
| C2—C3—C4 | 113.47 (14) | C4—C4B—H43B | 109.5 |
| C2—C3—H3A | 108.9 | H41B—C4B—H43B | 109.5 |
| C4—C3—H3A | 108.9 | H42B—C4B—H43B | 109.5 |
| C2—C3—H3B | 108.9 | C9B—C9C—H91C | 109.5 |
| C4—C3—H3B | 108.9 | C9B—C9C—H92C | 109.5 |
| H3A—C3—H3B | 107.7 | H91C—C9C—H92C | 109.5 |
| C6—C5—C4 | 114.60 (16) | C9B—C9C—H93C | 109.5 |
| C6—C5—H5A | 108.6 | H91C—C9C—H93C | 109.5 |
| C4—C5—H5A | 108.6 | H92C—C9C—H93C | 109.5 |
| C6—C5—H5B | 108.6 | | |

| | | | |
|-----------------|--------------|-----------------|--------------|
| C6—C7—C2—N1 | 178.61 (17) | C10—C9—C9A—O9A | 173.47 (17) |
| C8—C7—C2—N1 | -3.0 (3) | C8—C9—C9A—O9A | -8.8 (2) |
| C6—C7—C2—C3 | 0.7 (3) | C7—C2—C3—C4 | -26.3 (3) |
| C8—C7—C2—C3 | 179.07 (17) | N1—C2—C3—C4 | 155.73 (17) |
| C10—N1—C2—C7 | -11.5 (3) | C4B—C4—C3—C2 | -70.4 (2) |
| C10—N1—C2—C3 | 166.61 (17) | C4A—C4—C3—C2 | 169.08 (18) |
| C2—C7—C8—C9 | 17.0 (2) | C5—C4—C3—C2 | 49.6 (2) |
| C6—C7—C8—C9 | -164.56 (16) | O6A—C6—C5—C4 | -150.90 (19) |
| C2—C7—C8—C8A | -107.0 (2) | C7—C6—C5—C4 | 30.1 (3) |
| C6—C7—C8—C8A | 71.4 (2) | C3—C4—C5—C6 | -52.3 (2) |
| C7—C8—C9—C10 | -19.7 (2) | C4B—C4—C5—C6 | 67.8 (2) |
| C8A—C8—C9—C10 | 105.08 (19) | C4A—C4—C5—C6 | -172.20 (18) |
| C7—C8—C9—C9A | 162.58 (16) | C8C—C8B—C8A—C8F | -0.2 (3) |
| C8A—C8—C9—C9A | -72.7 (2) | C8C—C8B—C8A—C8 | 179.03 (19) |
| C2—C7—C6—O6A | 178.46 (18) | C7—C8—C8A—C8B | 55.7 (2) |
| C8—C7—C6—O6A | 0.0 (3) | C9—C8—C8A—C8B | -68.1 (2) |
| C2—C7—C6—C5 | -2.5 (3) | C7—C8—C8A—C8F | -125.0 (2) |
| C8—C7—C6—C5 | 179.01 (17) | C9—C8—C8A—C8F | 111.2 (2) |
| C9A—C9—C10—N1 | -174.18 (17) | C8B—C8A—C8F—C8E | 1.0 (3) |
| C8—C9—C10—N1 | 8.1 (3) | C8—C8A—C8F—C8E | -178.3 (2) |
| C9A—C9—C10—C10A | 4.3 (3) | C8A—C8B—C8C—O8C | -179.0 (2) |
| C8—C9—C10—C10A | -173.38 (18) | C8A—C8B—C8C—C8D | -0.2 (4) |
| C2—N1—C10—C9 | 8.7 (3) | C9A—O9A—C9B—C9C | -176.61 (19) |
| C2—N1—C10—C10A | -169.93 (18) | C8A—C8F—C8E—C8D | -1.3 (4) |
| C9B—O9A—C9A—O9B | -4.7 (3) | C8F—C8E—C8D—C8C | 0.8 (4) |
| C9B—O9A—C9A—C9 | 175.31 (17) | O8C—C8C—C8D—C8E | 178.7 (2) |
| C10—C9—C9A—O9B | -6.5 (3) | C8B—C8C—C8D—C8E | 0.0 (4) |
| C8—C9—C9A—O9B | 171.30 (19) | | |

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

| $D-H\cdots A$ | $D-H$ | $H\cdots A$ | $D\cdots A$ | $D-H\cdots A$ |
|-----------------------------------|-------|-------------|-------------|---------------|
| O8C—H8C \cdots O9B ⁱ | 0.82 | 2.05 | 2.835 (2) | 162 |
| N1—H1 \cdots O6A ⁱⁱ | 0.86 | 2.16 | 2.970 (2) | 157 |

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

Fig. 1

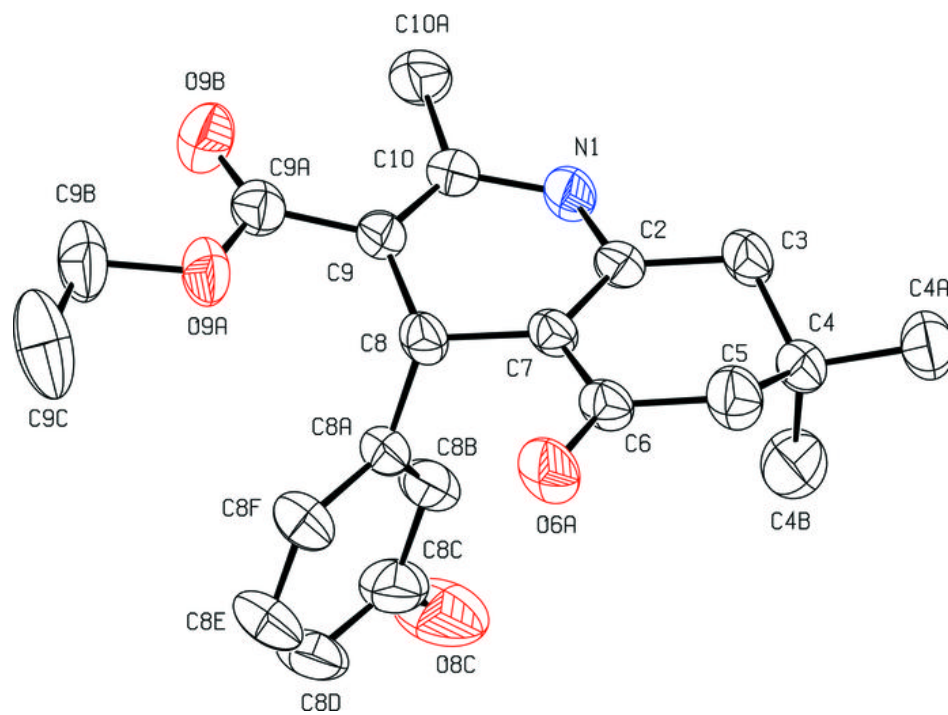


Fig. 2

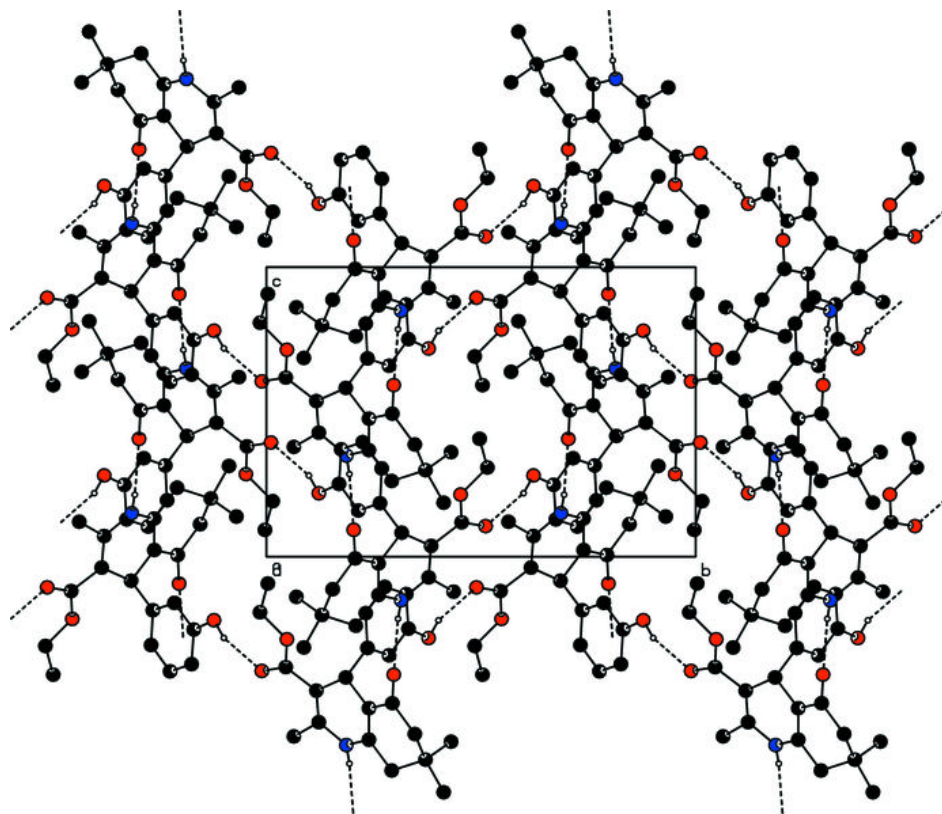


Fig. 3

