

# 3,3,6,6-Tetramethyl-9-(4-pyridyl)- 3,4,6,7,9,10-hexahydro-1,8(2*H*,5*H*)- acridinedione monohydrate

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## Key indicators

Single-crystal X-ray study

$T = 293\text{ K}$

Mean  $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$

$R$  factor = 0.051

$wR$  factor = 0.146

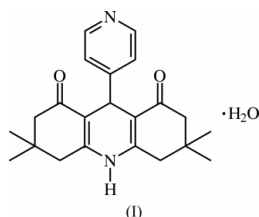
Data-to-parameter ratio = 18.5

For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

In the acridine moiety of the title compound,  $\text{C}_{22}\text{H}_{26}\text{N}_2\text{O}_2 \cdot \text{H}_2\text{O}$ , the central dihydropyridine ring adopts a flattened boat conformation, while the outer cyclohexene rings adopt sofa conformations. In the crystal structure,  $\text{N}-\text{H} \cdots \text{O}$  and  $\text{O}-\text{H} \cdots \text{O}$  hydrogen bonds involving the water molecule and  $\text{C}-\text{H} \cdots \text{O}$  hydrogen bonds link the inversion-related molecules to form layers parallel to the (011) plane. Adjacent layers are linked by  $\text{O}-\text{H} \cdots \text{N}$  hydrogen bonds involving the water molecule.

## Comment

Acridine derivatives exhibit a wide range of biological activities, especially mutagenic, antitumour (Talacki *et al.*, 1974) and anti-amoebic activities (Prasad Krishna *et al.*, 1984). Acridine-containing drugs have been found to possess anti-protozoal activity (Karolak-Wojciechowska *et al.*, 1996) and are used for the treatment of Alzheimer's disease (Bandoli *et al.*, 1994). The ability of acridine to intercalate between the base pairs of DNA is well known (Neidle, 1979; Fan *et al.*, 1997). Substituted hexahydroacridine-1,8-dione, which resembles K-channel openers, relaxes KCl-preconcentrated urinary-bladder smooth muscle *in vitro* (Li *et al.*, 1996; Trivedi *et al.*, 1995). Acridine-1,8-diones exhibit fluorescence and laser activities (Selladurai *et al.*, 1990). Acridinediones were found to lase around 475–495 nm (Murugan *et al.*, 1998). The present study of the title compound, (I), is part of a series of investigations on the crystal structures of acridinedione derivatives.

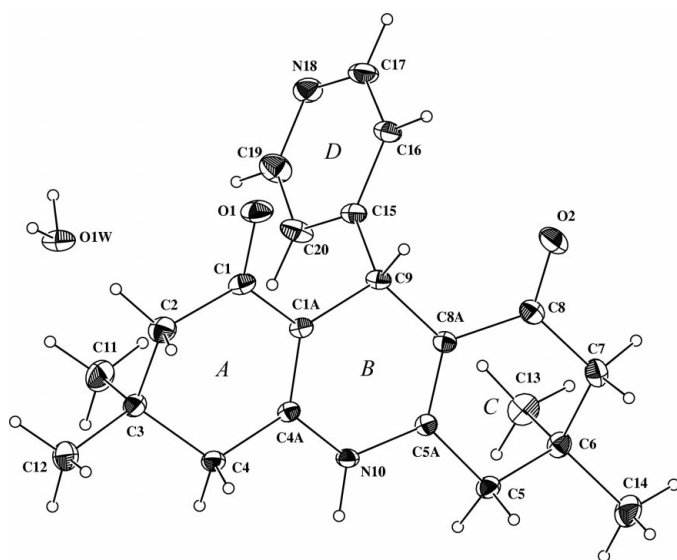


In the acridine moiety, the central pyridine ring (*B*) adopts a flattened boat conformation, while the outer rings (*A* and *C*) adopt sofa conformations as confirmed by the total puckering amplitudes (Cremer & Pople, 1975) [ $Q_T = 0.470$  (2), 0.144 (2) and 0.469 (2)  $\text{\AA}$  for rings *A*, *B* and *C*, respectively] and the asymmetry parameters (Nardelli, 1983) [ $\Delta_S(\text{C1A}) = 0.032$  (1),  $\Delta_S(\text{C9}) = 0.007$  (1) and  $\Delta_S(\text{C1A}-\text{C4A}) = 0.021$  (1),  $\Delta_S(\text{C6}) = 0.001$  (1) for rings *A*, *B* and *C*, respectively]. The puckering of ring *B* is quite small, owing to the  $\pi$  conjugation in the  $\text{C1A}-\text{C4A}-\text{N10}-\text{C5A}-\text{C8A}$  system, as indicated by the bond distances:  $\text{C1A}-\text{C4A} = 1.360$  (2),  $\text{C4A}-\text{N10} = 1.366$  (2),  $\text{N10}-\text{C5A} = 1.377$  (2) and  $\text{C5A}-\text{C8A} = 1.357$  (2)  $\text{\AA}$ . Similar features have also been observed in other acridinedione

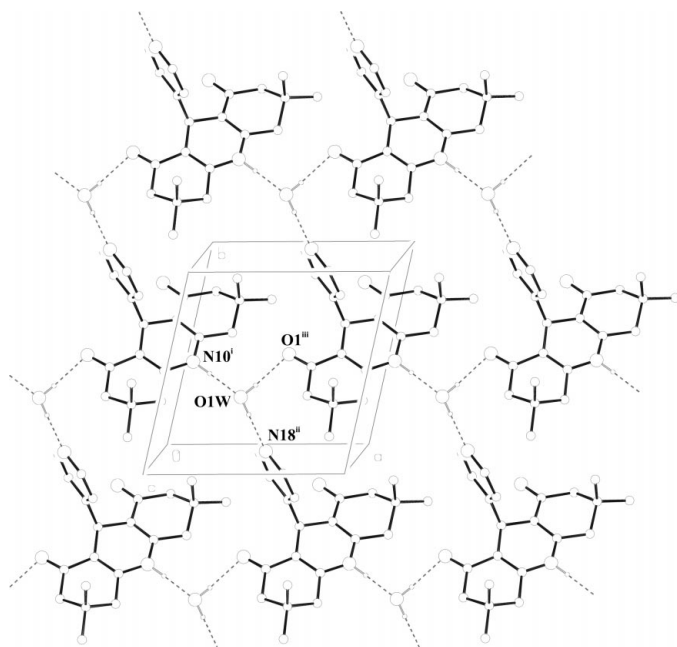
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**Figure 1**  
The molecular structure of title compound, showing 35% probability displacement ellipsoids.



**Figure 2**  
A view of the hydrogen-bonding network involving the water molecules [symmetry codes: (i)  $1-x, 1-y, 1-z$ ; (ii)  $1-x, -y, 1-z$ ; (iii)  $-x, 1-y, 1-z$ ]. For clarity, H atoms not involved in hydrogen bonding have been omitted.

analogues (Gunasekaran *et al.*, 1997; Ganesh, Banumathi *et al.*, 1998; Ganesh *et al.*, 1999; Sankaranarayanan *et al.*, 1998). The C4A–C1A–C9–C15 torsion angle of  $111.8(2)^\circ$  shows that the pyridyl ring (*D*) is axial to the acridine moiety. The acridine moiety is folded about the line passing through atoms C9 and N10, as seen from the dihedral angle of  $14.38(3)^\circ$  between the C1/C2/C4/C4A/N10/C9/C1A and C5/C7/C8/C8A/C9/N10/C5A planes. The folding of the acridine moiety about the C9···N10 line is well documented (Ganesh, Velmurugan *et al.*, 1998; Ganesh, Banumathi *et al.*, 1998; Ganesh *et al.*, 1999;

Sankaranarayanan *et al.*, 1998, 1999; Jeyakanthan *et al.*, 2000, 2002). The sum of the bond angles around N10 [ $359.3(1)^\circ$ ] indicates  $sp^2$  hybridization. In ring *B*, the C–N bond lengths (Table 1) are in agreement with the mean  $Csp^2-Nsp^2$  bond length of  $1.355(14)$  Å reported by Allen *et al.* (1987).

The hydrogen-bonding network involving the water molecules is shown in Fig. 2. The water molecules take part in O–H···O, N–H···O and O–H···N hydrogen bonds. The N10–H10···O1W<sup>i</sup>, O1W–H2W···O1<sup>iii</sup>, C11–H11B···O1<sup>iii</sup> and C17–H17···O2<sup>iv</sup> (symmetry codes as in Table 2) hydrogen bonds link the inversion-related molecules to form layers parallel to the (011) plane. Adjacent layers are linked by O1W–H1W···N18<sup>ii</sup> hydrogen bonds.

## Experimental

To dimedone (0.75 g, 5.3 mmol) and pyridine-4-carboxaldehyde (0.28 g, 2.6 mmol) in ethanol was added ammonia (excess) and the mixture was refluxed for 8 h to afford the title compound. Single crystals were grown by slow evaporation from a solution in chloroform–methanol (1:1).

### Crystal data

$C_{22}H_{26}N_2O_2 \cdot H_2O$   
 $M_r = 368.46$   
Triclinic,  $P\bar{1}$   
 $a = 9.1333(5)$  Å  
 $b = 9.8999(5)$  Å  
 $c = 12.0435(7)$  Å  
 $\alpha = 74.876(1)^\circ$   
 $\beta = 81.705(1)^\circ$   
 $\gamma = 73.137(1)^\circ$   
 $V = 1003.26(9)$  Å<sup>3</sup>

$Z = 2$   
 $D_x = 1.220$  Mg m<sup>-3</sup>  
Mo  $K\alpha$  radiation  
Cell parameters from 3984 reflections  
 $\theta = 1.8$ – $28.3^\circ$   
 $\mu = 0.08$  mm<sup>-1</sup>  
 $T = 293(2)$  K  
Plate, yellow  
 $0.48 \times 0.26 \times 0.12$  mm

### Data collection

Siemens SMART CCD area-detector diffractometer  
 $\omega$  scans  
7019 measured reflections  
4804 independent reflections  
3516 reflections with  $I > 2\sigma(I)$

$R_{int} = 0.022$   
 $\theta_{max} = 28.3^\circ$   
 $h = -12 \rightarrow 12$   
 $k = -13 \rightarrow 12$   
 $l = -13 \rightarrow 16$

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.051$   
 $wR(F^2) = 0.146$   
 $S = 1.03$   
4804 reflections  
260 parameters  
H atoms treated by a mixture of independent and constrained refinement

$w = 1/[\sigma^2(F_o^2) + (0.0728P)^2 + 0.1035P]$   
where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} = 0.001$   
 $\Delta\rho_{max} = 0.23$  e Å<sup>-3</sup>  
 $\Delta\rho_{min} = -0.19$  e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

O1–C1	1.2352 (18)	C9–C15	1.531 (2)
O2–C8	1.2255 (19)	C17–N18	1.333 (2)
C4A–N10	1.3660 (17)	N18–C19	1.328 (2)
C5A–N10	1.3774 (18)		
C4A–N10–C5A	121.49 (12)	C19–N18–C17	116.01 (15)
C1A–C9–C15–C20	−43.72 (19)	C8A–C9–C15–C16	−100.75 (16)
C1A–C9–C15–C16	135.57 (14)		

**Table 2**

Hydrogen-bonding geometry (Å, °).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N10-H10 \cdots O1W^i$	0.93 (2)	1.87 (2)	2.799 (2)	174 (2)
$O1W-H1W \cdots N18^{ii}$	0.91 (3)	1.96 (3)	2.869 (2)	179 (2)
$O1W-H2W \cdots O1^{iii}$	0.85 (3)	2.06 (3)	2.902 (2)	175 (2)
$C11-H11B \cdots O1^{iii}$	0.96	2.47	3.326 (2)	149
$C17-H17 \cdots O2^{iv}$	0.93	2.44	3.366 (2)	172

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

Atoms H10, H1W and H2W were located from a difference Fourier map and refined isotropically; the remaining H atoms were fixed geometrically and allowed to ride on their attached atoms. For the refined H atoms, the O–H distances are 0.85 (3) and 0.91 (3) Å, and the N–H distance is 0.93 (2) Å. Rotating-group refinement was used for the methyl groups.

Data collection: *SMART* (Siemens, 1996); cell refinement: *SAINT* (Siemens, 1996); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ZORTEP* (Zsolnai, 1997) and *PLATON* (Spek, 1990); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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