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#### Key indicators

Single-crystal X-ray study

$T = 120\text{ K}$

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

$R$  factor = 0.033

$wR$  factor = 0.085

Data-to-parameter ratio = 14.4

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

# Methyl ( $\pm$ )-(1 $\alpha$ ,2 $\beta$ ,8 $\alpha$ ,9 $\alpha$ ,10 $\beta$ )-2-chloro-4-aza-3-oxatetracyclo[8.4.0.0<sup>2,9</sup>.0<sup>4,8</sup>]-tetradecane-9-carboxylate oxalic acid monohydrate

In the crystal structure of the title compound,  $\text{C}_{14}\text{H}_{20}\text{ClNO}_3 \cdot \text{C}_2\text{H}_2\text{O}_4 \cdot \text{H}_2\text{O}$ , there are infinite flat ribbons, parallel to the  $[0\bar{1}1]$  direction. Molecules in the ribbons are linked together by an extensive network of hydrogen bonds.

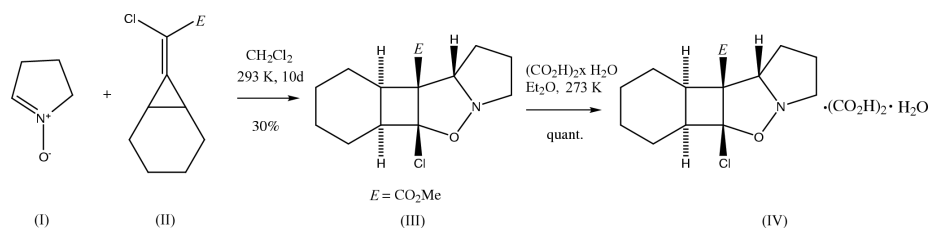
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## Comment

The tetracyclic title compound, (III), was obtained upon an attempted 1,3-dipolar cycloaddition of 3,4-dihydro-2*H*-pyrrole 1-oxide, (I), to methyl 2-chloro-2-(bicyclo[4.1.0]hept-7-ylidene)acetate, (II) (Brandi *et al.*, 2003), during systematic studies of 1,3-dipolar cycloadditions to substituted methyl-enecyclopropanes (Goti *et al.*, 1996; de Meijere, Kozhushkov & Khlebnikov, 2000; de Meijere & Kozhushkov, 2000). The relative configuration of (III) was established by an X-ray crystal structure analysis of its solvate with oxalic acid and water (1:1:1) (IV).



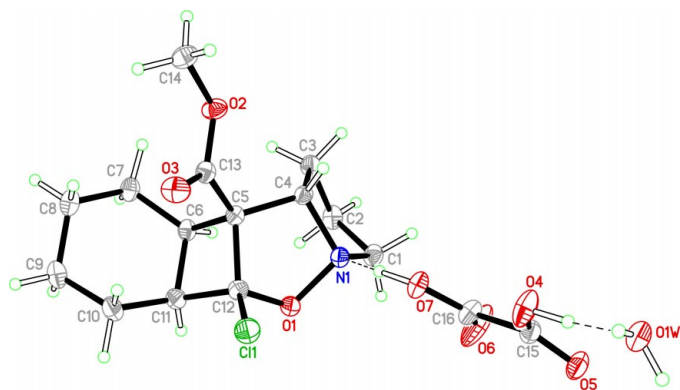
Chlorine and methoxycarbonyl substituents at the two bridgehead atoms of the tetracyclic system are *cis*-oriented with respect to each other (Fig. 1).

All four rings are substantially non-planar. The four-membered ring is folded around the  $\text{C6}\cdots\text{C12}$  line by  $24.7(1)^\circ$ . The cyclohexane ring has a slightly flattened chair conformation, while the five-membered heterocycles adopt conformations of O1-envelope and twisted half-chair. The solvent molecule of oxalic acid is also not entirely planar, with a torsion angle  $\text{O5}-\text{C15}-\text{C16}-\text{O6}$  of  $-7.5(2)^\circ$ .

The most remarkable feature is the packing of the molecules in the crystal. The molecules of oxalic acid and water form flat ribbons; the tetracyclic molecules are attached to both edges of the ribbons by very strong hydrogen bonds between oxalic acid and the tetradecane N atom (Fig. 2). The molecules of oxalic acid and water are linked together by a network of  $\text{O}-\text{H}\cdots\text{O}$  hydrogen bonds. Tetracyclic molecules in the same ribbon are also connected by  $\text{C6}-\text{H6}\cdots\text{O3}$  [ $\text{C}\cdots\text{O} = 3.290(1)\text{ \AA}$ ] interactions. In the crystal structure, these ribbons are parallel to the  $[0\bar{1}1]$  direction and show a number of weak  $\text{C}-\text{H}\cdots\text{O}$  interactions.

## Experimental

Crystals of (IV) were obtained by slow evaporation from a solution in  $\text{Et}_2\text{O}$  at 277 K.



**Figure 1**  
View of the asymmetric unit. Displacement ellipsoids are shown at the 50% probability level.

#### Crystal data

$C_{14}H_{20}ClNO_3 \cdot C_2H_2O_4 \cdot H_2O$   
 $M_r = 393.81$   
 Triclinic,  $P\bar{1}$   
 $a = 6.6133$  (2) Å  
 $b = 8.0737$  (2) Å  
 $c = 17.8848$  (5) Å  
 $\alpha = 77.746$  (1)°  
 $\beta = 83.211$  (1)°  
 $\gamma = 80.169$  (1)°  
 $V = 916.14$  (4) Å<sup>3</sup>

$Z = 2$   
 $D_x = 1.428$  Mg m<sup>-3</sup>  
 Mo  $K\alpha$  radiation  
 Cell parameters from 999 reflections  
 $\theta = 10.3$ – $20.8$ °  
 $\mu = 0.25$  mm<sup>-1</sup>  
 $T = 120$  (2) K  
 Block, colourless  
 $0.42 \times 0.35 \times 0.17$  mm

#### Data collection

Bruker SMART-CCD 1K diffractometer  
 $\omega$  scans  
 Absorption correction: none  
 11202 measured reflections  
 4763 independent reflections

3835 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.026$   
 $\theta_{max} = 29.0$ °  
 $h = -8 \rightarrow 9$   
 $k = -10 \rightarrow 10$   
 $l = -24 \rightarrow 23$

#### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.033$   
 $wR(F^2) = 0.085$   
 $S = 1.02$   
 4763 reflections  
 331 parameters  
 All H-atom parameters refined

$w = 1/[\sigma^2(F_o^2) + (0.04P)^2 + 0.3P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} < 0.001$   
 $\Delta\rho_{max} = 0.39$  e Å<sup>-3</sup>  
 $\Delta\rho_{min} = -0.28$  e Å<sup>-3</sup>

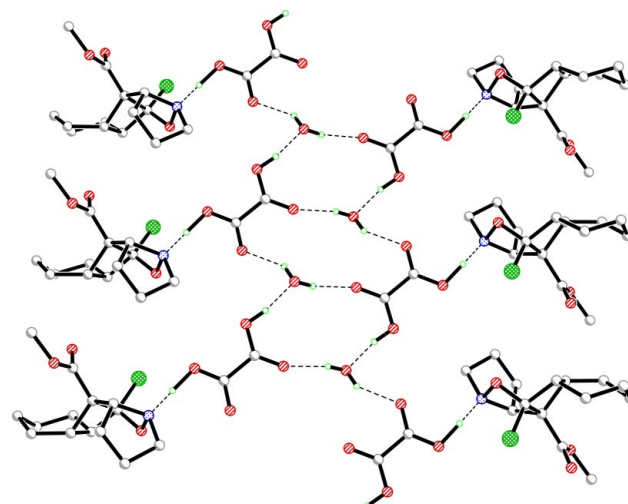
**Table 1**

Hydrogen-bonding geometry (Å, °).

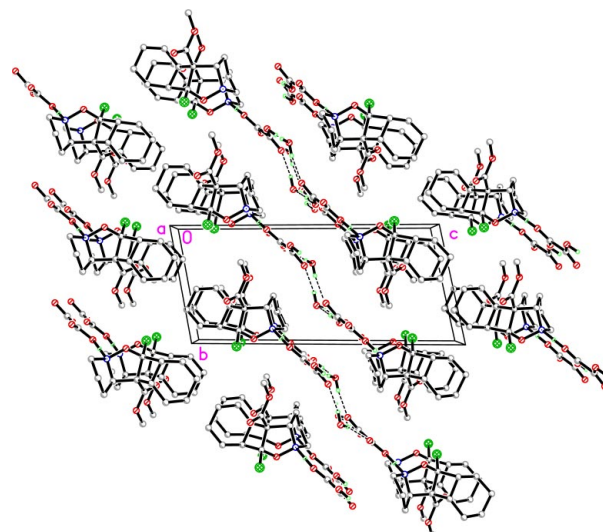
$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
O7–H7O <sup>i</sup> ···N1	1.22 (2)	1.34 (2)	2.5690 (14)	179 (2)
O4–H4O <sup>i</sup> ···O1W	0.94 (3)	1.63 (3)	2.5589 (14)	169 (2)
O1W–H1W <sup>i</sup> ···O5 <sup>i</sup>	0.86 (2)	1.94 (2)	2.7917 (14)	168 (2)
O1W–H2W <sup>i</sup> ···O6 <sup>ii</sup>	0.83 (2)	1.95 (2)	2.6886 (15)	147 (2)

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

Data collection: *SMART* (Bruker, 1997); cell refinement: *SMART*; data reduction: *SHELXTL* (Bruker, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics:



**Figure 2**  
Arrangement of the molecules in hydrogen-bonded ribbons.



**Figure 3**  
Packing diagram, viewed along the  $a$  axis.

*SHELXTL*; software used to prepare material for publication: *SHELXTL*.

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