

## 10,11-Dihydrocarbamazepine–acetic acid (1/1)

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## Key indicators

Single-crystal X-ray study  
 $T = 123$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.005$  Å  
 $R$  factor = 0.076  
 $wR$  factor = 0.147  
Data-to-parameter ratio = 12.5For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

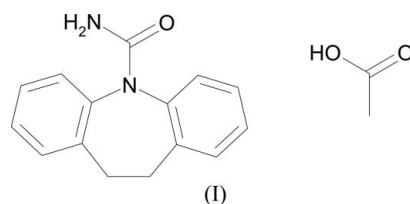
In the title compound [systematic name: 10,11-dihydro-5*H*-dibenz[*b,f*]azepine-5-carboxamide–ethanoic acid (1/1)],  $\text{C}_{15}\text{H}_{14}\text{N}_2\text{O}\cdot\text{C}_2\text{H}_4\text{O}_2$ , the dihydrocarbamazepine and acetic acid molecules are hydrogen bonded to form an  $R_2^2(8)$  motif, which is further connected into a centrosymmetric double motif arrangement.

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## Comment

10,11-Dihydrocarbamazepine (DHC) is a recognized impurity in carbamazepine (CBZ), a dibenzazepine drug used to control seizures (Cyr *et al.*, 1987). DHC is known to crystallize in three polymorphic forms: monoclinic form I (Bandoli *et al.*, 1992), orthorhombic form II (Harrison *et al.*, 2006) and triclinic form III (Leech *et al.*, 2006). The title compound, (I), was produced during an automated parallel crystallization study (Florence, Johnston, Fernandes *et al.*, 2006) of DHC as part of a wider study into the predicted and experimental structures of CBZ (Florence, Johnston, Price *et al.*, 2006; Florence, Leech *et al.*, 2006). The sample was identified as a new form using multi-sample foil transmission X-ray powder diffraction analysis (Florence *et al.*, 2003). Subsequent manual recrystallization from a saturated acetic acid solution by slow evaporation at 298 K yielded single crystals of (I) suitable for X-ray diffraction.



The crystal structure of (I) is essentially isostructural with that of CBZ–acetic acid (1/1) (Fleischman *et al.*, 2003). Accordingly, it displays the same space group with very similar unit-cell parameters and packing arrangements. Specifically, the DHC and acetic acid molecules are connected *via*  $\text{O2}-\text{H1}\cdots\text{O1}$  and  $\text{N2}-\text{H2N}\cdots\text{O3}$  hydrogen bonds (Table 1) to form an  $R_2^2(8)$  (Etter, 1990) dimer motif (Fig. 1). A third hydrogen bond,  $\text{N2}-\text{H1N}\cdots\text{O3}^i$  [symmetry code (i)  $1-x, 1-y, -z$ ], joins adjacent dimers to form a centrosymmetric double motif arrangement (Fig. 2).

## Experimental

Crystals of (I) were grown from a saturated acetic acid solution of 10,11-dihydrocarbamazepine by isothermal solvent evaporation at 298 K.

Crystal data

$C_{15}H_{14}N_2O \cdot C_2H_4O_2$   
 $M_r = 298.33$   
 Monoclinic,  $P2_1/c$   
 $a = 5.3104 (4) \text{ \AA}$   
 $b = 15.4246 (17) \text{ \AA}$   
 $c = 18.732 (2) \text{ \AA}$   
 $\beta = 95.106 (7)^\circ$   
 $V = 1528.3 (3) \text{ \AA}^3$

$Z = 4$   
 $D_x = 1.297 \text{ Mg m}^{-3}$   
 Mo  $K\alpha$  radiation  
 $\mu = 0.09 \text{ mm}^{-1}$   
 $T = 123 (2) \text{ K}$   
 Needle, colourless  
 $0.35 \times 0.08 \times 0.04 \text{ mm}$

Data collection

Nonius KappaCCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: none  
 10078 measured reflections

2652 independent reflections  
 1605 reflections with  $I > 2\sigma(I)$   
 $R_{int} = 0.103$   
 $\theta_{max} = 25.0^\circ$

Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.076$   
 $wR(F^2) = 0.147$   
 $S = 1.13$   
 2652 reflections  
 212 parameters

$w = 1/[\sigma^2(F_o^2) + (0.0421P)^2 + 0.9028P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} < 0.001$   
 $\Delta\rho_{max} = 0.20 \text{ e \AA}^{-3}$   
 $\Delta\rho_{min} = -0.21 \text{ e \AA}^{-3}$

H atoms treated by a mixture of independent and constrained refinement

Table 1

Hydrogen-bond geometry ( $\text{\AA}, ^\circ$ ).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O2-H1 \cdots O1$	1.03 (4)	1.53 (4)	2.547 (3)	167 (4)
$N2-H1N \cdots O3^i$	0.88 (4)	2.20 (3)	2.894 (4)	136 (3)
$N2-H2N \cdots O3$	0.95 (4)	2.04 (4)	2.970 (4)	164 (4)

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

H atoms bonded to N and O were located in difference maps and refined isotropically (distances are given in Table 1). All other H atoms were positioned geometrically and treated as riding with  $C-H = 0.95-0.99 \text{ \AA}$ , and with  $U_{iso}(H) = 1.2U_{eq}(C)$ , or  $U_{iso}(H) = 1.5U_{eq}(C)$  for the methyl group.

Data collection: COLLECT (Hooft, 1988) and DENZO (Otwinowski & Minor, 1997); cell refinement: DENZO and COLLECT; data reduction: DENZO; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: PLATON (Spek, 2003); software used to prepare material for publication: SHELXL97.

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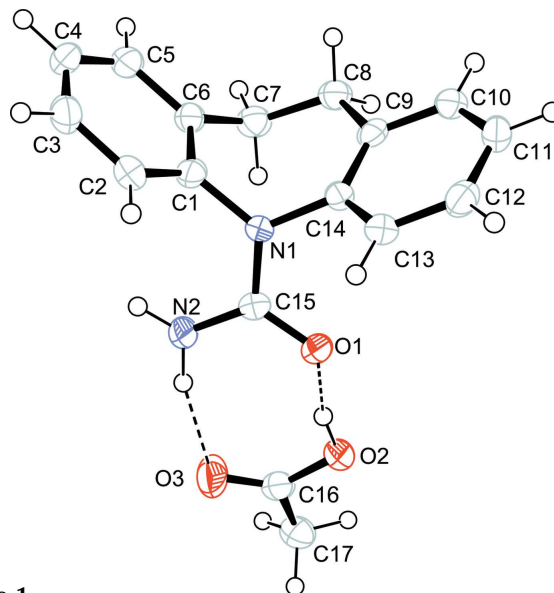


Figure 1 The asymmetric unit of (I), showing 50% probability displacement ellipsoids. Hydrogen bonds are shown as dashed lines.

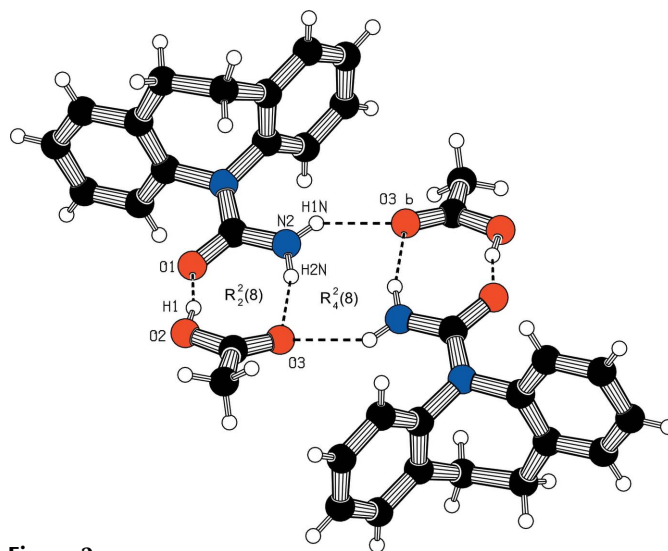


Figure 2 The hydrogen bonded  $R_2^2(8)$  motifs of (I) joined in a centrosymmetric arrangement via an  $R_4^4(8)$  motif. Hydrogen bonds are shown as dashed lines. [Symmetry code: (b)  $1 - x, 1 - y, -z$ .]

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