# organic compounds

Acta Crystallographica Section C Crystal Structure Communications ISSN 0108-2701

# A comparison of the supramolecular structures of 1-(6-amino-1,3-benzodioxol-5-yl)-3-(3,4,5-trimethoxyphenyl)prop-2-en-1-one and 1-(6-amino-1,3-benzodioxol-5-yl)-3-[4-(*N*,*N*-dimethylamino)phenyl]prop-2-en-1-one

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Received 20 August 2001 Accepted 30 October 2001 Online 22 December 2001

1-(6-Amino-1,3-benzodioxol-5-yl)-3-(3,4,5-trimethoxyphenyl)prop-2-en-1-one,  $C_{19}H_{19}NO_6$ , (I), contains an intramolecular  $N-H\cdots O$  hydrogen bond and a weak  $C-H\cdots O$  hydrogen bond, which forms a C(10) chain motif running parallel to the c axis. The 6-amino-1,3-benzodioxol-5-yl moieties are involved in  $\pi-\pi$  stacking. 1-(6-Amino-1,3-benzodioxol-5-yl)-3-[4-(N,Ndimethylamino)phenyl]prop-2-en-1-one,  $C_{18}H_{18}N_2O_3$ , (II), crystallizes with two molecules in the asymmetric unit. The main feature of the supramolecular structure of (II) is the formation of a centrosymmetric hydrogen-bonded tetramer with an  $R_8^4(16)$  motif.

## Comment

Many synthetic or naturally occurring compounds containing the 1,3-dioxolyl group are very important because of their pharmacological properties (Ma *et al.*, 1987; Ohta & Kimoto, 1976; Schlunke & Egli, 1972; Krause & Goeber, 1972; Gabrielsen *et al.*, 1992). Chalcones (1,3-diarylpropenones) have been widely used as starting materials in numerous synthetic reactions (Awad *et al.*, 1960; Coudert *et al.*, 1988; Carrie & Rochard, 1963), including the preparation of fusedring heterocyclic compounds (Insuasty *et al.*, 1992, 1997; Kolos *et al.*, 1996). We have prepared the title compounds, (I) and (II), as intermediates in the synthesis of novel dihydroquinolinones. This is a class of compounds with interesting pharmaceutical properties (Pfizer, 1980; Jordis *et al.*, 1991; Baker *et al.*, 1990), since the 1,3-dioxolyl residue in their frameworks enhances their practical applications.



The molecular structures of (I) and the two independent molecules of (II) are shown in Figs. 1 and 2, respectively. The



### Figure 1

A view of the molecule of (I) with the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.



#### Figure 2

Views of the two independent molecules of (II) with the atom-numbering schemes. Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radii.

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dioxol-5-yl groups in all three molecules have C2 envelope puckers. In (I), a strong intramolecular hydrogen bond is observed with an S(6) motif via N5-H5A···O8 (Bernstein et



#### Figure 3

A view of the crystal structure of (I), looking down the *b* axis, showing two anti-parallel *C*(10) chains [symmetry code: (i) x,  $-\frac{1}{2} - y$ ,  $z - \frac{1}{2}$ ].





A view of the tetramer formed by the molecules of (II) at (x, y, z) and (1 - x, 1 - y, 1 - z). For the sake of clarity, H atoms bonded to C atoms have been omitted [symmetry code: (ii) 1 - x, 1 - y, 1 - z].



The supramolecular structures of (I) and (II) are completely different. In (I), a C(10) motif is formed via C16-H16···O1(x,  $-\frac{1}{2} - y$ ,  $z - \frac{1}{2}$ ), creating a chain which runs parallel to the c axis. Two such anti-parallel chains are shown in Fig. 3. Atom H5B is not involved in any hydrogen bonding. In (II), however, the equivalent H atoms H15B and H25B are involved in strong hydrogen bonds via N15- $N15B \cdots O28(1 - x, 1 - y, 1 - z)$  and  $N25 - H25B \cdots O18$ , linking molecules 1 and 2 to form a tetramer with an  $R_8^4(16)$ motif (Fig. 4). Such motifs occur centred at  $(l + \frac{1}{2}, m + \frac{1}{2}, n + \frac{1}{2})$ , where l, m and n are integers or zero. The weak C14- $H14 \cdots O28(1 - x, 1 - y, 1 - z)$  and  $C24 - H24 \cdots O18$ hydrogen bonds help to reinforce this tetramer structure. The tetramers are linked to each other by weak hydrogen bonds via C212-H212···O13(x, y, z - 1) and C212(1 - x, 1 - y, z)-H212(1 - x, 1 - y, z)···O13(1 - x, 1 - y, 1 - z), which form bridging  $R_4^4(26)$  rings along with the N15-H15...O28 hydrogen bonds of the tetramer (Fig. 5).

In (I), there is  $\pi$ - $\pi$  stacking between the benzo-1,3-dioxol-5-yl groups, where the rings defined by O1/C2/O3/C3a/C7a and C3a/C4-C7/C7a are stacked such that the distance between the centres of gravity of the rings is 3.7276 (9) Å, the dihedral angle is 3.50°, the perpendicular distance from the centre of gravity of the former ring at (x, y, z) to the latter at (x, 1 + y, z) is 3.368 Å and the perpendicular distance from the centre of gravity of the latter ring at (x, y, z) to the former at



## Figure 5

A view of the  $R_4^4(26)$  rings which link the tetramers in (II) [symmetry codes: (ii) 1 - x, 1 - y, 1 - z; (iii) x, y, z - 1; (iv) 1 - x, 1 - y, -z]. For the sake of clarity, H atoms bonded to C or N atoms and not involved in this motif have been omitted, as has the unit-cell box.

(x, y - 1, z) is 3.445 Å (Fig. 6). It is worth noting that the dihedral angle between these planes, 3.50°, is in fact the dihedral angle between the two components of the benzo-1,3dioxol-5-yl group, which is not in itself planar. There is no  $\pi - \pi$ stacking in (II).



## Figure 6

A view of the  $\pi \cdots \pi$  stacking in (I). For the sake of clarity the unit-cell box has been omitted.

## **Experimental**

For (I), a mixture of 6'-amino-3',4'-(methylenedioxy)acetophenone (0.50 g, 2.79 mmol), 3,4,5-trimethoxybenzaldehyde (2.79 mmol), ethanol (10 ml) and NaOH (0.5 ml, 20% aq.) was heated to 333-343 K for 10-15 min and the reaction monitored by thin-layer chromatography. After cooling, the solid which formed was filtered off and washed with ethanol to yield pure (I) as orange crystals (93% yield; m.p. 464 K). For (II), a mixture of 6'-amino-3',4'-(methylenedioxy)acetophenone (0.50 g, 2.79 mmol), 4-(N,N-dimethylamino)benzaldehyde (2.79 mmol), ethanol (10 ml) and NaOH (0.5 ml, 20% aq.) was heated to 333-343 K for 10-15 min and the reaction monitored by thin-layer chromatography. After cooling, the solid which formed was filtered off and washed with ethanol to yield pure (II) as purple crystals (46% yield; m.p. 435 K). For (I) and (II), crystals suitable for single-crystal X-ray diffraction were selected directly from the prepared samples.

## Compound (I)

## Crystal data

$C_{19}H_{19}NO_{6}$	$D_x = 1.451 \text{ Mg m}^{-3}$
$M_r = 357.35$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 3622
$a = 23.5477 (4) \text{\AA}$	reflections
b = 4.9605 (1)  Å	$\theta = 3.1-27.5^{\circ}$
c = 14.2556 (8) Å	$\mu = 0.11 \text{ mm}^{-1}$
$\beta = 100.702 \ (1)^{\circ}$	T = 150 (1)  K
$V = 1636.21 (10) \text{ Å}^3$	Block, orange
Z = 4	$0.4 \times 0.1 \times 0.1 \text{ mm}$

#### Data collection

Nonius KappaCCD diffractometer $\varphi$ and $\omega$ scans with $\kappa$ offsets Absorption correction: multi-scan ( <i>DENZO-SMN</i> ; Otwinowski & Minor, 1997) $T_{min} = 0.958, T_{max} = 0.989$ 11 152 measured reflections	3622 independent reflections 2832 reflections with $I > 2\sigma(I)$ $R_{int} = 0.048$ $\theta_{max} = 27.5^{\circ}$ $h = -30 \rightarrow 30$ $k = -6 \rightarrow 6$ $l = -18 \rightarrow 18$
Refinement	
Refinement on $F^2$ $R[F^2 > 2\sigma(F^2)] = 0.044$ $wR(F^2) = 0.124$ S = 1.03 3622 reflections 238 parameters H-atom parameters constrained	$w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0675P)^{2} + 0.3552P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3 (\Delta/\sigma)_{\text{max}} < 0.001$ $\Delta\rho_{\text{max}} = 0.26 \text{ e} \text{ Å}^{-3} \Delta\rho_{\text{min}} = -0.32 \text{ e} \text{ Å}^{-3}$

#### Table 1

Hydrogen-bonding geometry (Å,  $^{\circ}$ ) for (I).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D{\cdots}A$	$D - H \cdots A$
$\begin{array}{c} N5-H5A\cdotsO8\\ C16-H16\cdotsO1^{i}\end{array}$	0.88	2.01	2.651 (2)	129
	0.95	2.51	3.398 (2)	155

Symmetry code: (i)  $x, \frac{1}{2} - y, \frac{1}{2} + z$ .

### Compound (II)

Crystal data	
$C_{18}H_{18}N_2O_3$	Z = 4
$M_r = 310.34$	$D_x = 1.339 \text{ Mg m}^{-3}$
Triclinic, P1	Mo $K\alpha$ radiation
a = 9.5849(2)  Å	Cell parameters from 6884
b = 12.3218 (4) Å	reflections
c = 13.3304 (4) Å	$\theta = 3.1 - 27.5^{\circ}$
$\alpha = 83.4617 (14)^{\circ}$	$\mu = 0.09 \text{ mm}^{-1}$
$\beta = 80.2411 (17)^{\circ}$	T = 150 (1)  K
$\gamma = 85.7271 (16)^{\circ}$	Block, purple
$V = 1539.04 (8) Å^3$	$0.22 \times 0.14 \times 0.12 \text{ mm}$
Data collection	
Nonius KappaCCD	6884 independent reflections
diffractometer	4873 reflections with $I > 2\sigma(I)$
$\varphi$ and $\omega$ scans with $\kappa$ offsets	$R_{\rm int} = 0.085$
Absorption correction: multi-scan	$\theta_{\rm max} = 27.5^{\circ}$
(DENZO-SMN; Otwinowski &	$h = -12 \rightarrow 12$
Minor, 1997)	$k = -15 \rightarrow 15$
$T_{\min} = 0.980, \ T_{\max} = 0.989$	$l = -17 \rightarrow 17$
23 078 measured reflections	

## Table 2

Hydrogen-bonding geometry (Å,  $^{\circ}$ ) for (II).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
N15-H15A···O18	0.88	2.03	2.6433 (14)	125
N15-H15 $B$ ···O28 <sup>i</sup>	0.88	2.26	3.0867 (13)	157
N25-H25A···O28	0.88	1.94	2.6237 (17)	133
$N25 - H25B \cdots O18$	0.88	2.20	3.0606 (17)	165
$C14-H14\cdots O28^{i}$	0.95	2.38	3.2067 (18)	145
C24-H24···O18	0.95	2.52	3.3212 (19)	142
$C212{-}H212{\cdot}{\cdot}{\cdot}O13^{ii}$	0.95	2.60	3.4508 (18)	149

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

Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.054$	$w = 1/[\sigma^2 (F_o^2) + (0.0927P)^2]$
$wR(F^2) = 0.154$	where $P = (F_o^2 + 2F_c^2)/3$
S = 1.02	$(\Delta/\sigma)_{\rm max} = 0.006$
6884 reflections	$\Delta \rho_{\rm max} = 0.30 \text{ e } \text{\AA}^{-3}$
419 parameters	$\Delta \rho_{\rm min} = -0.34 \ {\rm e} \ {\rm \AA}^{-3}$

For both compounds, H atoms were treated as riding, with C–H = 0.95-0.99 Å. In the case of the N–H bonds, the H atoms were initially located on a difference Fourier map. The bond lengths were then *DFIX*ed (*SHELXL97*; Sheldrick, 1997) to 0.88 Å with an s.u. of 0.005 until a value close to 0.88 Å was obtained, at which stage they were *AFIX*ed (*SHELXL97*) and allowed to ride on their parent atoms.

For both compounds, data collection: *KappaCCD Server Software* (Nonius, 1997); cell refinement: *DENZO-SMN* (Otwinowski & Minor, 1997); data reduction: *DENZO-SMN*; program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEP*II (Johnson, 1976) and *PLATON* (Spek, 2000); software used to prepare material for publication: *SHELXL*97 and *PRPKAPPA* (Ferguson, 1999).

X-ray data were collected at the EPSRC X-ray Crystallographic Service at the University of Southampton and the authors thank the staff there for all their help and advice. The authors also thank COLCIENCIAS and Universidad del Valle for financial support.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: GD1178). Services for accessing these data are described at the back of the journal.

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