

4-Ethynyl-1,2-methylenedioxybenzene  
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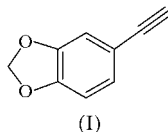
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The title compound, C<sub>9</sub>H<sub>6</sub>O<sub>2</sub>, contains two moderate C—H···O hydrogen bonds. That involving the terminal alkyne gives rise to chains along the *b* axis. The other hydrogen bond occurs over a centre of symmetry, leading to dimers. The combination of the two interactions gives rise to rings, each comprising six molecules, which are part of infinite sheets in the *bc* plane.

## Comment

The packing of simple aromatic compounds has attracted interest (Desiraju & Kishan, 1989; Weiss *et al.*, 1997). It has been demonstrated by Sarma & Desiraju (1986) that the effects of heteroatoms (principally O or Cl) either within the aromatic ring or peripheral to it can be structure determining. A low number of heteroatoms means that structures tend to adopt herring-bone motifs; benzene, with none, is the archetypal example (Bacon *et al.*, 1964). On the other hand, higher numbers of heteroatoms can lead to stacked motifs, *e.g.* pyranthrene (Sarma & Desiraju, 1986). Desiraju & Kishan (1989) demonstrated this principle with structural studies on 3,4-(methylenedioxy)cinnamic acid and [3,4-(methylenedioxy)phenyl]propionic acid. Therefore, we decided to examine the structural motifs present in the related compound 4-ethynyl-1,2-methylenedioxybenzene, (I).



Compound (I) lies in a general position (Fig. 1), but with the non-H atoms coplanar to within 0.008 Å. All molecular geometry parameters lie within the normal ranges (Allen *et al.*, 1987).

The most significant feature of (I) is the presence of two intermolecular C—H···O hydrogen bonds (Table 1). The first occurs over a centre of symmetry so that the aromatic H2 atom acts as a donor to O9<sup>ii</sup> [symmetry code: (ii) 1 - *x*, 1 - *y*,

-*z*], generating a dimeric centrosymmetric R<sub>2</sub><sup>2</sup>(8) motif (Fig. 2).

The second intermolecular hydrogen bond involves the alkynyl H8 atom, which acts as a donor to *n*-glide-related O9<sup>i</sup> [symmetry code: (i)  $\frac{3}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$ ]. This generates C(7) chains along the *y* axis, leading to herring-bone chains in the *ab* plane. The top half of each dimer is related to the bottom half of the neighbouring dimer by this C—H···O interaction (Fig. 2). Therefore, the dimers are related to each other in a zigzag manner along the *b* direction within the *bc* plane. The

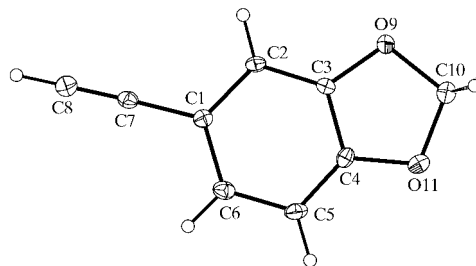


Figure 1

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

combination of the two hydrogen bonds generates a six-molecule R<sub>6</sub><sup>4</sup>(28) ring as part of an infinite two-dimensional sheet (Fig. 2). There is also a possible C5—H5···O11<sup>iii</sup> interaction [symmetry code: (iii) -*x*, 1 - *y*, 1 - *z*] between dimers related by a centre of symmetry in the middle of the R<sub>6</sub><sup>4</sup>(28) ring.

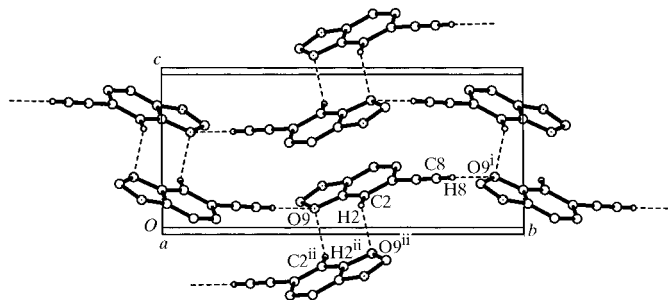


Figure 2

The packing diagram of (I) showing part of an infinite two-dimensional sheet in the *bc* plane. The component dimeric units and six-molecule rings are visible [symmetry codes: (i)  $\frac{3}{2} - x, \frac{1}{2} + y, \frac{1}{2} - z$ ; (ii) 1 - *x*, 1 - *y*, -*z*].

## Experimental

Compound (I) (Pelter *et al.*, 1990) was prepared according to the method of Corey & Fuchs (1972). 3,4-(Methylenedioxy)benzaldehyde was firstly transformed to the dibromoalkene by treatment with carbon tetrabromide, triphenylphosphine and zinc; subsequent treatment with butyllithium generated the title alkyne, (I), which exists as a supercooled oil at room temperature. The sample was placed in a sealed capillary and cooled *in situ* on the diffractometer using an Oxford Cryosystems open-flow cryostat (Cosier & Glazer, 1986). The crystal was grown by repeated cycling of the temperature across the freezing point of the compound (260 K).

## Crystal data

$C_9H_6O_2$	$D_x = 1.411 \text{ Mg m}^{-3}$
$M_r = 146.14$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/n$	Cell parameters from 40 reflections
$a = 6.318 (3) \text{ \AA}$	$\theta = 10.2\text{--}12.6^\circ$
$b = 15.867 (10) \text{ \AA}$	$\mu = 0.100 \text{ mm}^{-1}$
$c = 7.571 (4) \text{ \AA}$	$T = 150 (2) \text{ K}$
$\beta = 115.03 (3)^\circ$	Needle, pale yellow
$V = 687.7 (7) \text{ \AA}^3$	$0.19 \times 0.11 \times 0.08 \text{ mm}$
$Z = 4$	

## Data collection

Stoe Stadi-4 four-circle diffractometer	$\theta_{\text{max}} = 26.04^\circ$
$\omega/\theta$ scans	$h = -7 \rightarrow 7$
5424 measured reflections	$k = -19 \rightarrow 19$
1356 independent reflections	$l = -9 \rightarrow 9$
729 reflections with $I > 2\sigma(I)$	3 standard reflections
$R_{\text{int}} = 0.196$	frequency: 60 min
	intensity decay: 9.6%

## Refinement

Refinement on $F^2$	H-atom parameters constrained
$R[F^2 > 2\sigma(F^2)] = 0.086$	$w = 1/[\sigma^2(F_o^2) + 1.6470P]$
$wR(F^2) = 0.163$	where $P = (F_o^2 + 2F_c^2)/3$
$S = 1.153$	$(\Delta/\sigma)_{\text{max}} < 0.001$
1356 reflections	$\Delta\rho_{\text{max}} = 0.31 \text{ e \AA}^{-3}$
100 parameters	$\Delta\rho_{\text{min}} = -0.35 \text{ e \AA}^{-3}$

Table 1

Geometry of hydrogen bonds and short intermolecular contacts ( $\text{\AA}$ ,  $^\circ$ ).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$C8-H8 \cdots O9^i$	0.95	2.58	3.526 (7)	178
$C2-H2 \cdots O9^{ii}$	0.95	2.53	3.344 (7)	143
$C5-H5 \cdots O11^{iii}$	0.95	2.70	3.529 (7)	146

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

All H atoms were clearly defined in difference maps and were then treated as riding, with C—H = 0.95 or 0.99  $\text{\AA}$ . Examination of the structure with *PLATON* (Spek, 2000) showed that there were no

solvent-accessible voids in the crystal lattice. The high value of  $R_{\text{int}}$  is attributed to the presence, within the region of the capillary irradiated by the X-ray beam, of additional crystallites of (I), the presence of which is an unavoidable consequence of the crystal-growth method.

Data collection: *STADIA* (Stoe & Cie, 1997); cell refinement: *STADIA*; data reduction: *X-RED* (Stoe & Cie, 1997); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2000); software used to prepare material for publication: *SHELXL97* and *PLATON*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: GD1130). Services for accessing these data are described at the back of the journal.

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