

## (E)-1-[4-(Hexyloxy)phenyl]-3-(2-hydroxy-phenyl)prop-2-en-1-one

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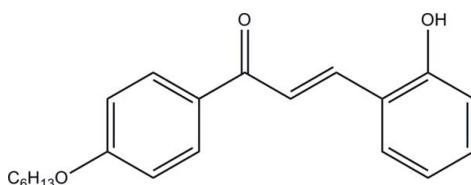
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Key indicators: single-crystal X-ray study;  $T = 100$  K; mean  $\sigma(C-C) = 0.002 \text{ \AA}$ ;  $R$  factor = 0.041;  $wR$  factor = 0.133; data-to-parameter ratio = 20.6.

In the title compound,  $C_{21}H_{24}O_3$ , the enone moiety adopts an *s-cis* conformation and the dihedral angle between the benzene rings is  $12.89(6)^\circ$ . The hexyloxy tail adopts an extended conformation. In the crystal, inversion dimers are linked by pairs of  $O-H\cdots O$  hydrogen bonds and pairs of  $C-H\cdots O$  interactions, forming two  $R_2^2(7)$  and one  $R_2^2(10)$  loops. The dimers are then arranged into sheets lying parallel to (011) and weak  $C-H\cdots\pi$  interactions consolidate the packing.

### Related literature

For a related structure and background to the biological properties of chalcones, see: Ngaini *et al.* (2011). For related structures, see: Razak *et al.* (2009); Ngaini *et al.* (2010). For graph-set theory, see: Bernstein *et al.* (1995). For the stability of the temperature controller used in the data collection, see: Cosier & Glazer (1986).



### Experimental

#### Crystal data

$C_{21}H_{24}O_3$   
 $M_r = 324.40$   
Triclinic,  $P\bar{1}$

$a = 7.485(2) \text{ \AA}$   
 $b = 10.834(3) \text{ \AA}$   
 $c = 11.673(3) \text{ \AA}$

‡ Thomson Reuters ResearcherID: A-5599-2009.

$\alpha = 73.858(5)^\circ$   
 $\beta = 77.961(6)^\circ$   
 $\gamma = 76.941(6)^\circ$   
 $V = 874.9(4) \text{ \AA}^3$   
 $Z = 2$

Mo  $K\alpha$  radiation  
 $\mu = 0.08 \text{ mm}^{-1}$   
 $T = 100 \text{ K}$   
 $0.47 \times 0.14 \times 0.12 \text{ mm}$

#### Data collection

Bruker APEX DUO CCD diffractometer  
Absorption correction: multi-scan (*SADABS*; Bruker, 2009)  
 $T_{\min} = 0.963$ ,  $T_{\max} = 0.991$

17565 measured reflections  
4576 independent reflections  
3781 reflections with  $I > 2\sigma(I)$   
 $R_{\text{int}} = 0.026$

#### Refinement

$R[F^2 > 2\sigma(F^2)] = 0.041$   
 $wR(F^2) = 0.133$   
 $S = 1.02$   
4576 reflections  
222 parameters

H atoms treated by a mixture of independent and constrained refinement  
 $\Delta\rho_{\max} = 0.39 \text{ e \AA}^{-3}$   
 $\Delta\rho_{\min} = -0.23 \text{ e \AA}^{-3}$

**Table 1**  
Hydrogen-bond geometry ( $\text{\AA}$ ,  $^\circ$ ).

$Cg1$  is the centroid of the C1–C6 ring.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O2-H1O2\cdots O1^i$	0.94 (2)	1.78 (2)	2.6862 (15)	163.3 (19)
$C7-H7A\cdots O2^i$	0.93	2.37	3.2526 (18)	158
$C16-H16A\cdots Cg1^{ii}$	0.97	2.91	3.6117 (16)	130

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

Data collection: *APEX2* (Bruker, 2009); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXTL* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL*; molecular graphics: *SHELXTL*; software used to prepare material for publication: *SHELXTL* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB6948).

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