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## Crystal Structure

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# threo-2-(2,6-Dimethoxyphenoxy)-1-(4-hydroxy-3,5-dimethoxyphenyl)-propane-1,3-diol: a conformational study 

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In the crystal structure of the title compound, $\mathrm{C}_{19} \mathrm{H}_{24} \mathrm{O}_{8}$, the molecules adopt a conformation in which the bulky 2,6 -dimethoxyphenoxy and 4-hydroxy-3,5-dimethoxyphenyl groups are almost as far apart as possible. The C(aryl)…C(aryl) distance is 4.8766 (19) $\AA$, which is close to the calculated maximum value ( $4.92 \AA$ ). The $\mathrm{C}($ aryl) $-\mathrm{C}-\mathrm{C}-\mathrm{O}$ (aryloxy) torsion angle is $173.76(11)^{\circ}$ and the C (benzylic) $-\mathrm{C}-\mathrm{O}-$ C (aryl) torsion angle is $149.09(11)^{\circ}$. The conformation is compared with those of related lignin model compounds. The hydrogen-bonding pattern is discussed in terms of graph-set theory.

## Comment

This paper describes the crystal structure of the title compound, (I), the threo form of a lignin model compound representative of structural elements in lignin of the arylglycerol $\beta$-syringyl ether type. The crystal structure of the erythro form, (II), has been reported previously (Langer, $\mathrm{Li} \&$ Lundquist, 2002). Arylglycerol $\beta$-syringyl ethers constitute a major type of structural elements in hardwood lignins, e.g. in birch lignin (Larsson \& Miksche, 1971). The erythro form of such structural elements predominates (Lundquist \& von Unge, 1986; Bardet et al., 1998; Akiyama et al., 2003).

In (I), the $\mathrm{C} 9-\mathrm{C} 10-\mathrm{O} 6-\mathrm{C} 12$ torsion angle is $149.09(11)^{\circ}$ and the $\mathrm{C} 1-\mathrm{C} 9-\mathrm{C} 10-\mathrm{O} 6$ torsion angle is $173.76(11)^{\circ}$ (the enantiomer with the $R$ configuration at the benzylic C atom is considered throughout the discussion of torsion angles in this paper), and the $\mathrm{C} 1 \cdots \mathrm{C} 12$ distance is $4.8766(19) \AA$. In the erythro form, the corresponding data are $-75.26(13)^{\circ}$, $-177.27(10)^{\circ}$ and 4.4458 (17) Å, respectively. The C1 . . C12 distance in (II) is about the same as the corresponding distances in other examined erythro forms of the arylglycerol $\beta$-syringyl ether type (Stomberg \& Lundquist, 1989; Langer \&

Lundquist, 2001; Langer et al., 2005). Obviously, the aromatic rings in the erythro form, (II) (and related compounds), are not separated as much as they are in the threo form, (I). The angle between the aromatic ring planes is $67.66(6)^{\circ}$ in (I) and 57.27 (5) ${ }^{\circ}$ in (II). The C1 . . C12 distance in (I) [4.8766 (19) A $]$ is very close to the calculated maximum value ( $4.92 \AA$ ).

(I)

(III) $R=\mathrm{H}, R^{\prime}=\mathrm{McO}$
(IV) $R=R^{\prime}=\mathrm{H}$
(V) $R=\mathrm{Mc}, R^{\prime}=\mathrm{H}$

In the crystal structure of (I), there are three intramolecular (Fig. 1) and two intermolecular hydrogen bonds of the O $\mathrm{H} \cdots \mathrm{O}$ type and four weak intermolecular hydrogen bonds of the $\mathrm{C}-\mathrm{H} \cdots \mathrm{O}$ type (Table 1). On the first-level graph-set (Bernstein et al., 1995; Grell et al., 1999), the intramolecular hydrogen bonds are classified as $S(5)$ for bonds $a$ and $c$, and $S(8)$ for $d$ (see Fig. 1). The intermolecular hydrogen bonds $b$ and $e$ form $C(10)$ and $C(6)$ chains, respectively (Fig. 2). The weak intramolecular hydrogen bonds $f, g, h$ and $i$ form $R_{2}^{2}(8)$ rings and $C(8), C(13)$ and $C(7)$ chains, respectively. On the second-level graph-set, many chains and rings could be identified, the most important ones being $C_{2}^{2}(10)$ and $C_{2}^{2}(16)$ chain types, both formed by bonds $b$ and $e$. These two hydrogen bonds thus form $R_{3}^{3}(22)$ rings (Fig. 2). The assignment of graph-set descriptors was performed using PLUTO, as described by Motherwell et al. (1999). The hydrogen-bonding patterns of threo-(I) and erythro-(II) (Langer, Li \& Lundquist, 2002) forms are similar with respect to intramolecular hydrogen bonds.

As pointed out above, the title threo $\beta$-syringyl ether adopts a conformation in which the aryl groups are almost as far apart as possible. This is actually what could be expected from computational studies (Besombes et al., 2003a). In a second threo form of an arylglycerol $\beta$-syringyl ether, namely threo-2-(2,6-dimethoxyphenoxy)-1-(3,4-dimethoxyphenyl)propane-1,3-diol, the $\mathrm{C}($ aryl $)-\mathrm{C}-\mathrm{C}-\mathrm{O}$ (aryloxy) torsion angle is -70.5 (2) ${ }^{\circ}$ and the $\mathrm{C}($ aryl $)-\mathrm{O}-\mathrm{C}-\mathrm{C}($ benzylic) torsion angle is $-148.25(19)^{\circ}$ (Langer, Lundquist et al., 2002). This leads to a C(aryl)…C(aryl) distance of 4.319 (3) $\AA$, which deviates


Figure 1
A perspective drawing of (I), showing the atom-numbering scheme. Displacement ellipsoids are shown at the $50 \%$ probability level. The intramolecular hydrogen bonds are shown as dashed lines and their notation is given in Table 1.


Figure 2
The hydrogen-bonding pattern (dashed lines) of (I) in projection along the $c$ axis. The hydrogen-bond notation is given in Table 1.
significantly from the maximum value ( ca $4.91 \AA$ ). However, according to computational studies, the conformation adopted by this compound is fairly favoured (Besombes et al., 2003a). All the threo forms of models of the arylglycerol $\beta$-guaiacyl ether type which have been examined to date [(III) (Stomberg et al., 1988), (IV) and its triacetate (Lundquist et al., 1996), and (V) (Langer \& Lundquist, 2002)] adopt conformations in which the aromatic groups are far apart from each other. This is in accordance with expectations based on computational studies (Simon \& Eriksson, 1998; Besombes et al., 2003b). The crystal structure of the threo form of a trimeric model compound of the arylglycerol $\beta$-aryl ether type, (VI), has been reported by Karhunen et al. (1996). Calculations based on their data show that this compound also adopts an extended conformation: the $\mathrm{C}($ aryl $)-\mathrm{C}-\mathrm{C}-\mathrm{O}$ (aryloxy) torsion angle is $175.7^{\circ}$ and the $\mathrm{C}($ aryl $)-\mathrm{O}-\mathrm{C}-\mathrm{C}$ (benzylic) torsion angle is $171.9^{\circ}$.

## Experimental

Compound (I) was synthesized according to the procedure of Li et al. (2000). Separation of the erythro and threo forms was accomplished by ion-exchange chromatography (cf. Li et al., 1994). Crystals of (I) (m.p. 415-417 K) suitable for X-ray crystallography were obtained from 2-butanone/ethyl acetate (3:1).

Crystal data
$\mathrm{C}_{19} \mathrm{H}_{24} \mathrm{O}_{8}$
$M_{r}=380.38$
Monoclinic, C2/c
$a=24.0634$ (1) $\AA$
$b=7.3465$ (1) A
$c=22.4459$ (3) A
$\beta=114.332(1)^{\circ}$
$V=3615.56(7) \AA^{3}$
$Z=8$

## Data collection

Siemens SMART CCD areadetector diffractometer $\omega$ scans
Absorption correction: multi-scan (SADABS; Sheldrick, 2002)
$T_{\text {min }}=0.768, T_{\text {max }}=0.976$
28856 measured reflections

## Refinement

Refinement on $F^{2}$

$$
\begin{aligned}
& D_{x}=1.398 \mathrm{Mg} \mathrm{~m}^{-3} \\
& \text { Mo } K \alpha \text { radiation } \\
& \text { Cell parameters from } 7395 \\
& \quad \text { reflections } \\
& \theta=1.9-30.7^{\circ} \\
& \mu=0.11 \mathrm{~mm}^{-1} \\
& T=173(2) \mathrm{K} \\
& \text { Prism, colourless } \\
& 0.28 \times 0.24 \times 0.22 \mathrm{~mm}
\end{aligned}
$$

5566 independent reflections
4423 reflections with $I>2 \sigma(I)$
$R_{\text {int }}=0.035$
$\theta_{\text {max }}=30.7^{\circ}$
$h=-34 \rightarrow 34$
$k=-10 \rightarrow 10$
$l=-32 \rightarrow 32$
$R\left[F^{2}>2 \sigma\left(F^{2}\right)\right]=0.050$
$w R\left(F^{2}\right)=0.146$
$S=1.03$
5566 reflections
275 parameters
H atom parameters constrained

$$
\begin{aligned}
& w= 1 /\left[\sigma^{2}\left(F_{\mathrm{o}}^{2}\right)+(0.0693 P)^{2}\right. \\
&+4.0089 P] \\
& \text { where } P=\left(F_{\mathrm{o}}^{2}+2 F_{\mathrm{c}}^{2}\right) / 3 \\
&(\Delta / \sigma)_{\max }=0.001 \\
& \Delta \rho_{\max }=0.55 \mathrm{e} \AA^{-3} \\
& \Delta \rho_{\min }=-0.27 \mathrm{e}^{-3}
\end{aligned}
$$

H atoms were refined isotropically and were constrained to ideal geometry using an appropriate riding model. For aromatic H atoms, the $\mathrm{C}-\mathrm{H}$ distances were kept fixed at $0.95 \AA$, for secondary H atoms at $0.99 \AA$ and for tertiary H atoms at $1.00 \AA$. For the hydroxyl groups, the $\mathrm{O}-\mathrm{H}$ distance $\left(0.84 \AA\right.$ ) and $\mathrm{C}-\mathrm{O}-\mathrm{H}$ angle $\left(109.5^{\circ}\right)$ were kept fixed, while the torsion angle was allowed to refine, with the starting position based on a circular Fourier synthesis. For methyl groups, the $\mathrm{C}-\mathrm{H}$ distance $\left(0.98 \AA\right.$ ) and $\mathrm{C}-\mathrm{C}-\mathrm{H}$ angle $\left(109.5^{\circ}\right)$ were kept fixed, while the torsion angles were allowed to refine, with the starting positions based on a threefold averaged circular Fourier synthesis.

Table 1
Hydrogen-bonding geometry $\left(\AA,{ }^{\circ}\right)$.

| Label | $D-\mathrm{H} \cdots A$ | D-H | $\mathrm{H} \cdots A$ | $D \cdots A$ | $D-\mathrm{H} \cdots A$ |
| :---: | :---: | :---: | :---: | :---: | :---: |
| $a$ | $\mathrm{O} 2-\mathrm{H} 2 \cdots \mathrm{O} 3$ | 0.84 | 2.22 | 2.6707 (16) | 114 |
| $b$ | $\mathrm{O} 2-\mathrm{H} 2 \cdots \mathrm{O} 5^{\text {i }}$ | 0.84 | 2.02 | 2.7851 (15) | 152 |
| c | O4-H4 . . O6 | 0.84 | 2.31 | 2.7424 (14) | 112 |
| $d$ | O4-H4. . O7 | 0.84 | 2.13 | 2.9123 (16) | 154 |
| $e$ | $\mathrm{O} 5-\mathrm{H} 5 \cdots \mathrm{O} 4^{\text {ii }}$ | 0.84 | 1.99 | 2.8284 (15) | 173 |
| $f$ | C9-H9 . $\mathrm{O}^{\text {O }}{ }^{\text {iii }}$ | 1.00 | 2.56 | 3.5139 (17) | 159 |
| $g$ | $\mathrm{C} 11-\mathrm{H} 11 A \cdots \mathrm{O} 3^{\text {iv }}$ | 0.99 | 2.50 | 3.2171 (18) | 129 |
| $h$ | $\mathrm{C} 18-\mathrm{H} 18 \mathrm{C} \cdots \mathrm{O}^{\text {v }}$ | 0.98 | 2.57 | 3.466 (2) | 153 |
| , | $\mathrm{C} 19-\mathrm{H} 19 \mathrm{C} \cdots \mathrm{O}^{\text {ii }}$ | 0.98 | 2.52 | 3.491 (2) | 169 |

Symmetry codes: (i) $\frac{5}{2}-x, \frac{1}{2}+y, \frac{3}{2}-z$; (ii) $x, y-1, z ;$ (iii) $2-x, y, \frac{3}{2}-z$; (iv) $\frac{5}{2}-x, y-\frac{1}{2}$,
$\frac{3}{2}-z$; (v) $x-\frac{1}{2}, \frac{3}{2}-y, z-\frac{1}{2}$.

Data collection: SMART (Siemens, 1995); cell refinement: SAINT (Siemens, 1995); data reduction: SAINT and SADABS (Sheldrick, 2002); program(s) used to solve structure: SHELXTL (Bruker, 1997); program(s) used to refine structure: $S H E L X T L$; molecular graphics: DIAMOND (Brandenburg, 2005); software used to prepare material for publication: SHELXTL.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1820). Services for accessing these data are described at the back of the journal.

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