

(E)-2,4,5-Trimethoxybenzaldehyde oxime**Xin-Hong Chang**Department of Chemistry, Luoyang Teacher's
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The title compound, $C_{10}H_{13}NO_4$, which exists as the *E* isomer, crystallizes with two independent molecules in the asymmetric unit. The bond lengths and angles in both molecules are normal. The crystal structure is stabilized by intermolecular $O-H\cdots O$ hydrogen bonds, which link the molecules into cyclic tetramers.

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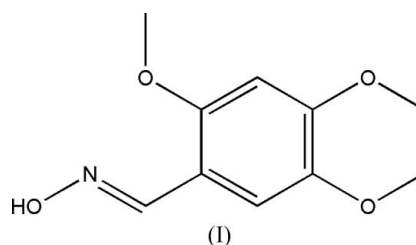
Key indicators

Single-crystal X-ray study
 $T = 293\text{ K}$
 Mean $\sigma(C-C) = 0.002\text{ \AA}$
 R factor = 0.035
 wR factor = 0.106
 Data-to-parameter ratio = 14.0

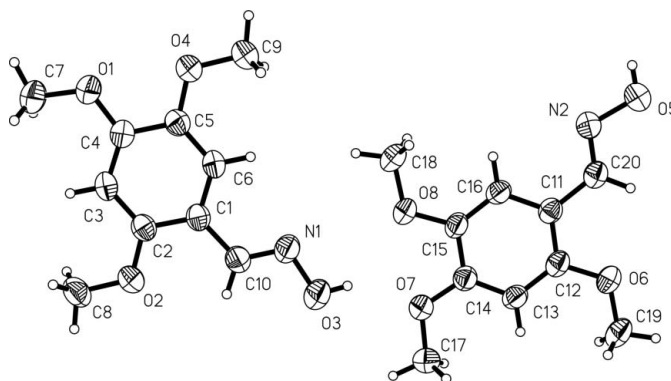
For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Comment

Substituted benzaldehyde oxime is an important intermediate in organic synthesis (Xu & Jin, 1999), existing in two isomeric forms, *z*. and *E* (Sharghi & Sarvari, 2001). We report here the crystal structure of the title compound, (I).



In (I) (Fig. 1), all bond lengths and angles are normal (Allen *et al.*, 1987) and in good agreement with those reported previously (Jerslev, 1983; Jensen, 1970). Atoms O1/O2/O4/C1–C9 lie in the same plane, the largest deviation being 0.040 (2) Å for atom C7, while in the second independent molecule, atoms C11–C20 and O6–O8 are essentially coplanar, the largest deviation from the mean plane being 0.046 (3) Å for atom C19. Intermolecular $O-H\cdots O$ hydrogen bonds (Table 2) link the molecules into nearly planar cyclic tetramers (Fig. 2).

**Figure 1**

The asymmetric unit of the title compound (I), with displacement ellipsoids drawn at the 40% probability level.

Experimental

The title compound was synthesized by the reaction of 2,4,5-trimethoxybenzaldehyde (0.01 mol) with hydroxylamine hydrochloride (0.01 mol) in the presence of sodium carbonate (0.01 mol) in a water–methanol solution (20 ml; 1:5 v/v) at room temperature (3.5 h). After dilution with water, the solution was extracted with dichloromethane, and the organic phase was evaporated to afford the title product in 92% isolated yield (1.94 g). Crystals suitable for X-ray diffraction analysis were obtained by slow evaporation of an ethyl acetate–petroleum ether (1:1 v/v) solution at room temperature over a period of two weeks.

Crystal data

$C_{10}H_{13}NO_4$ $V = 1081.04 (2) \text{ \AA}^3$
 $M_r = 211.21$ $Z = 4$
 Triclinic, $P\bar{1}$ $D_x = 1.298 \text{ Mg m}^{-3}$
 $a = 7.44280 (10) \text{ \AA}$ Mo $K\alpha$ radiation
 $b = 10.71890 (10) \text{ \AA}$ $\mu = 0.10 \text{ mm}^{-1}$
 $c = 14.0290 (2) \text{ \AA}$ $T = 293 (2) \text{ K}$
 $\alpha = 87.5830 (10)^\circ$ Block, colorless
 $\beta = 88.4510 (10)^\circ$ $0.36 \times 0.30 \times 0.22 \text{ mm}$
 $\gamma = 75.2140 (10)^\circ$

Data collection

Bruker SMART CCD area-detector 12599 measured reflections
 diffractometer 3805 independent reflections
 φ and ω scans 3079 reflections with $I > 2\sigma(I)$
 Absorption correction: multi-scan $R_{int} = 0.016$
 (SADABS; Sheldrick, 1996) $\theta_{max} = 25.0^\circ$
 $T_{min} = 0.965, T_{max} = 0.978$

Refinement

Refinement on F^2 $w = 1/[\sigma^2(F_o^2) + (0.0542P)^2 + 0.1378P]$
 $R[F^2 > 2\sigma(F^2)] = 0.035$ where $P = (F_o^2 + 2F_c^2)/3$
 $wR(F^2) = 0.106$ $(\Delta/\sigma)_{max} < 0.001$
 $S = 1.03$ $\Delta\rho_{max} = 0.15 \text{ e \AA}^{-3}$
 3805 reflections $\Delta\rho_{min} = -0.11 \text{ e \AA}^{-3}$
 272 parameters Extinction correction: SHELXTL
 H-atom parameters constrained Extinction coefficient: 0.0058 (13)

Table 1

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

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$O5-H5A \cdots O1^1$	0.82	2.14	2.8676 (15)	148
$O5-H5A \cdots O4^1$	0.82	2.29	2.9832 (15)	142

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

H atoms were placed in calculated positions, with $C-H = 0.93$ or 0.96 \AA and $O-H = 0.82 \text{ \AA}$, the hydroxy H atom being initially

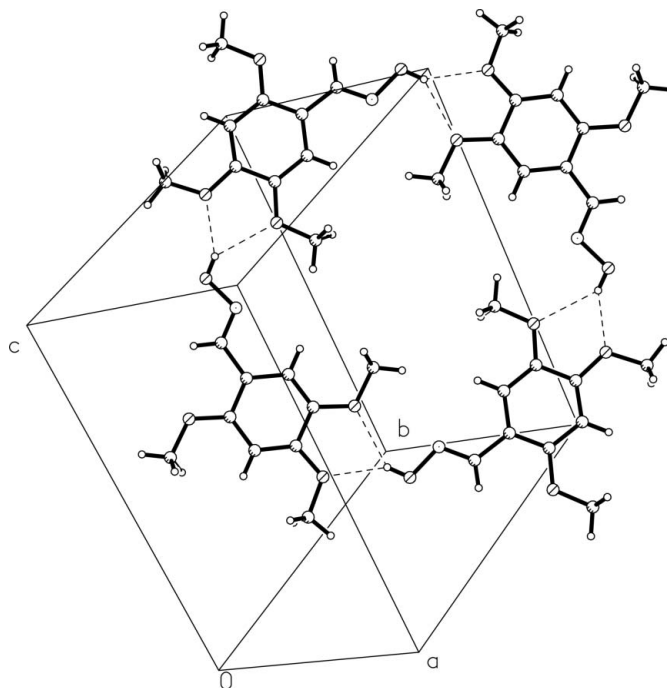


Figure 2

Partial packing plot, showing the hydrogen-bonded (dashed lines) tetramer.

located in a difference Fourier map. They were refined using a riding model, with $U_{iso}(H) = 1.2U_{eq}(C)$ for the aryl H atoms and $1.5U_{eq}(C,O)$ for the hydroxy and methyl H atoms.

Data collection: SMART (Bruker, 1998); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXTL (Bruker, 1999); program(s) used to refine structure: SHELXTL; molecular graphics: SHELXTL; software used to prepare material for publication: SHELXTL.

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