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#### **Key indicators**

Single-crystal X-ray study T = 298 K Mean  $\sigma$ (C–C) = 0.003 Å Disorder in main residue R factor = 0.054 wR factor = 0.148 Data-to-parameter ratio = 11.8

For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e. In the title compound,  $C_8H_6F_3NO$ , the trifluoromethyl group displays rotational disorder. The crystal packing is stabilized by intermolecular  $O-H\cdots N$  hydrogen bonds and van der Waals forces.

(E)-4-(Trifluoromethyl)benzaldehyde oxime

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## Comment

4-(Trifluoromethyl)benzaldehyde oxime is an important intermediate in organic synthesis, in particular in the formation of isoxazole (Sharghi & Sarvari, 2001). In this paper we report the crystal structure of the title compound, (I).



In (I), the bond lengths and angles are in agreement with values reported previously (Jerslev, 1983; Jensen, 1970). Atoms C1–C8, N1 and O1 are almost coplanar, the largest deviation being 0.050 (2) Å for atom N1. The crystal packing is stabilized by  $O-H\cdots N$  intermolecular hydrogen bonds (Table 1) and van der Waals forces.

## **Experimental**

The title compound was synthesized by the reaction of 4-(trifluoromethyl)benzaldehyde (0.01 mol) with hydroxylamine hydrochloride (0.01 mol) in the presence of sodium carbonate (0.01 mol) in an aqueous methanol (water: methanol 1:1  $\nu/\nu$ ) solution (20 ml) at room temperature (5 h). After diluting with water, the aqueous solution



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#### Figure 1

View of the title compound (I), with displacement ellipsoids drawn at the 40% probability level. Both disorder components are shown.

# organic papers

was extracted with dichloromethane, and the organic phase was dried and evaporated to afford the title product in 89% yield (Zhang & Li, 2004). Crystals suitable for X-ray diffraction analysis were obtained by slow evaporation of a solution in a hexane/dichloromethane mixture (1:1  $\nu/\nu$ ) at room temperature over a period of one week.

#### Crystal data

 $\begin{array}{l} C_8 H_6 F_3 \text{NO} \\ M_r = 189.14 \\ \text{Monoclinic, } P2_1/c \\ a = 7.543 \ (4) \ \text{\AA} \\ b = 7.395 \ (4) \ \text{\AA} \\ c = 15.058 \ (8) \ \text{\AA} \\ \beta = 99.278 \ (7)^{\circ} \\ V = 829.0 \ (8) \ \text{\AA}^3 \end{array}$ 

### Data collection

Bruker SMART CCD area-detector diffractometer  $\varphi$  and  $\omega$  scans Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)  $T_{\rm min} = 0.952, T_{\rm max} = 0.961$ 

#### Refinement

Refinement on  $F^2$   $R[F^2 > 2\sigma(F^2)] = 0.054$   $wR(F^2) = 0.148$  S = 1.061536 reflections 130 parameters H-atom parameters constrained Z = 4  $D_x = 1.515 \text{ Mg m}^{-3}$ Mo K\alpha radiation  $\mu = 0.15 \text{ mm}^{-1}$  T = 298 (2) KBlock, colorless  $0.36 \times 0.29 \times 0.24 \text{ mm}$ 

4169 measured reflections 1536 independent reflections 1259 reflections with  $I > 2\sigma(I)$  $R_{\text{int}} = 0.050$  $\theta_{\text{max}} = 25.5^{\circ}$ 

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.0725P)^{2} + 0.1821P]$ where  $P = (F_{o}^{2} + 2F_{c}^{2})/3$   $(\Delta/\sigma)_{max} < 0.001$   $\Delta\rho_{max} = 0.23 \text{ e} \text{ Å}^{-3}$   $\Delta\rho_{min} = -0.27 \text{ e} \text{ Å}^{-3}$ Extinction correction: SHELXL97
Extinction coefficient: 0.73 (5)

# Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$O1-H1\cdots N1^i$	0.82	2.12	2.831 (3)	145
Symmetry code: (i)	-x + 1, -y, -z.			

All H atoms were placed in calculated positions, with C-H = 0.93 Å and O-H = 0.82 Å, and included in the final cycles of refinement using a riding model, with  $U_{iso}(H) = 1.2U_{eq}(C)$  for the aryl H atoms and  $1.5U_{eq}(O)$  for the hydroxyl H atom. The trifluoromethyl group was found to be disordered; atoms C8, F1, F2 and F3 were refined over two positions [occupancies 0.093 (3) for the primed and 0.907 (3) for the unprimed atoms].

Data collection: *SMART* (Bruker, 1998); cell refinement: *SAINT* (Bruker, 1999); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *SHELXTL* (Bruker, 1999); software used to prepare material for publication: *SHELXTL*.

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