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2-Fluoro-4-(methoxycarbonyl)benzoic acid

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Key indicators: single-crystal X-ray study; T = 296 K; mean σ (C–C) = 0.004 Å; R factor = 0.066; wR factor = 0.190; data-to-parameter ratio = 12.0.

In the crystal of the title compound, $C_9H_7FO_4$, classical carboxylate inversion dimers are linked by pairs of $O-H\cdots O$ hydrogen bonds. The packing is consolidated by $C-H\cdots F$ and $C-H\cdots O$ interactions. The benzene ring and the methoxycarbonyl group are nearly coplanar, with a dihedral angle of 1.5 (3)° between them, whereas the carboxyl group has a dihedral angle of 20.2 (4)° with respect to the benzene ring.

Related literature

For background to the applications of the title compound, see: Jiang *et al.* (2008); Sakaki *et al.* (2007). For related structures, see: Wagner *et al.* (2009).

Experimental

Crystal data

 $C_9H_7FO_4$ $M_r = 198.15$

Triclinic, $P\overline{1}$ a = 7.536 (7) Å b = 7.591 (7) Å Z = 2 c = 8.523 (8) Å Mo Kα radiation α = 99.480 (14)° μ = 0.13 mm⁻¹ β = 108.748 (13)° T = 296 K γ = 99.240 (14)° $0.25 \times 0.19 \times 0.08$ mm V = 443.3 (7) Å³

Data collection

 $\begin{array}{ll} \mbox{Bruker SMART APEX CCD} & 2526 \mbox{ measured reflections} \\ \mbox{diffractometer} & 1535 \mbox{ independent reflections} \\ \mbox{Absorption correction: multi-scan} & 1025 \mbox{ reflections with } I > 2\sigma(I) \\ \mbox{} (SADABS; \mbox{ Bruker}, 2008) & R_{\rm int} = 0.025 \\ \mbox{} T_{\rm min} = 0.969, \ T_{\rm max} = 0.990 \\ \end{array}$

Refinement

 $\begin{array}{ll} R[F^2 > 2\sigma(F^2)] = 0.066 & 128 \ {\rm parameters} \\ WR(F^2) = 0.190 & {\rm H-atom\ parameters\ constrained} \\ S = 1.02 & \Delta\rho_{\rm max} = 0.24\ {\rm e\ \mathring{A}^{-3}} \\ 1535\ {\rm reflections} & \Delta\rho_{\rm min} = -0.22\ {\rm e\ \mathring{A}^{-3}} \end{array}$

Table 1Hydrogen-bond geometry (Å, °).

D $ H$ $\cdot \cdot \cdot A$	D-H	$H \cdot \cdot \cdot A$	$D \cdot \cdot \cdot A$	D $ H$ $\cdot \cdot \cdot A$
$ \begin{array}{c} C9-H9A\cdots F1^{i} \\ O2^{ii}-H2A^{ii}\cdots O1 \\ C3-H3A\cdots O3^{iii} \end{array} $	0.96 0.82 0.93	2.54 1.86 2.53	3.278 (5) 2.672 (4) 3.325 (4)	134 (1) 170 (1) 144 (1)
Symmetry codes: $-x$, $-y + 2$, $-z + 1$.	(i) $x - 1, y -$	-1, z-1; (ii)	-x+2, -y+3	-z+1; (iii)

Data collection: *APEX2* (Bruker, 2008); cell refinement: *SAINT-Plus* (Bruker, 2008); data reduction: *SAINT-Plus*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XSHELL* (Bruker, 2004); software used to prepare material for publication: *APEX2*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: PB2036).

References

Bruker (2004). XSHELL. Bruker AXS Inc., Madison, Wisconsin, USA Bruker (2008). APEX2, SADABS and SAINT-Plus. Bruker AXS Inc., Madison, Wisconsin, USA.

Jiang, X., Lee, G. T., Prasad, K. & Repic, O. (2008). Org. Process Res. Dev. 12, 1137–1141.

Sakaki, J., Kishida, M., Konishi, K., Gunji, H., Toyao, A., Matsumoto, Y., Kanazawa, T., Uchiyama, H., Fukaya, H., Mitani, H., Arai, Y. & Kimura, M. (2007). Bioorg. Med. Chem. Lett. 17, 4804–4807.

Sheldrick, G. M. (2008). Acta Cryst. A64, 112-122.

Wagner, C. E. et al. (2009). J. Med. Chem. 52, 5950-5966.

supplementary m	aterials	

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2-Fluoro-4-(methoxycarbonyl)benzoic acid

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Comment

The title compound, 4-(methoxycarbonyl)-2-fluorobenzoic acid, has recently been used to prepare novel diazepinylbenzoic acid retinoid-X-receptor antagonists (Jiang *et al.*, 2008; Sakaki *et al.*, 2007) as potential oral anti-obesity and anti-diabetic treatments as well as novel retinoid-X-receptor agonists with potential to treat various human cancers. Thus, the X-ray diffraction data of the present study confirms the fluorine locus for 4-(methoxycarbonyl)-2-fluorobenzoic acid.

The structure consists of sheets parallel to $(2\overline{1}2)$ stabilized by six intermolecular hydrogen interactions per molecule as shown in Table 1. The benzene ring and the methoxycarbonyl group are essentially coplanar as shown by the 1.51 (25)° dihedral angle between the two planes. However, the carboxylic acid is not coplanar with the benzene ring, as shown by the 20.18 (36)° dihedral angle between those two planes.

Experimental

The method of Sakaki and co-workers (Sakaki *et al.*, 2007) was followed to synthesize (1). To a flask containing 3-fluoro-4-formylmethylbenzoate (Wagner *et al.*, 2009) (9.22 g, 50.5 mmol) and sulfamic acid (5.40 g, 55.6 mmol) in water (21 ml) and ACN (42 ml) was slowly added a solution of 80% NaClO₂ (4.92 g, 53.8 mmol) in water (21 ml) at room temperature. After being stirred for 1 h, the reaction solution was added to a saturated, aqueous solution of Na₂SO₃ (75 ml) and 1 N HCl (150 ml), and the resulting solution was extracted with ethyl acetate (75 ml) three times. The combined organic extracts were washed with brine, dried over sodium sulfate, and the solvents were removed *in vacuo* to give crude (1) (7.56 g, 75%) as a white solid. A small sample was crystallized from hot ethyl acetate to give pure (1) as white crystals, m.p. 154–155 °C: 1 H NMR (400 MHz, CDCl₃) δ 10.5 (br s, 1H), 8.10 (t, J = 7.8, 1H), 7.89 (d, J = 8.2, 1H), 7.82 (d, J = 11.0, 1H), 3.97 (s, 3H); 13 C NMR (100.6 MHz, CDCl₃) δ 168.6, 168.5, 165.0, 164.9, 163.4, 160.8, 136.7, 136.6, 132.8, 124.9, 124.8, 121.3, 121.2, 118,4, 118.1, 52.8; LC-APCI-MS (M+) calcd for C₉H₇O₄F 198.0328, found 198.0331.

Refinement

H atoms were placed geometrically and allowed to refine as atoms riding on their bonding partners. The hydrogen was placed on the carboxylic acid based on the longer of the carboxylic acid carbon-oxygen bonds.

Figures

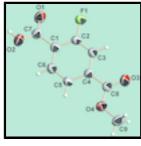


Fig. 1. Labeled thermal ellipsoid plot of (1) shown at the 50% probability level for all non-H atoms.

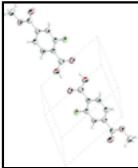


Fig. 2. Molecular pair of (1) shown at the 50% probability level for all non-H atoms illustrating classical intermolecular centrosymmetric carboxylic acid hydrogen bonding interactions.



Fig. 3. Packing diagram of (1) shown at the 50% probability level for all non-H atoms showing the alternating molecular orientations in two adjacent layers.

2-Fluoro-4-(methoxycarbonyl)benzoic acid

Crystal data

Z = 2C₉H₇FO₄ $M_r = 198.15$ F(000) = 204Triclinic, $P\overline{1}$ $D_{\rm x} = 1.484 \; {\rm Mg \; m}^{-3}$ Hall symbol: -P 1 Melting point: 427 K a = 7.536 (7) ÅMo $K\alpha$ radiation, $\lambda = 0.71073 \text{ Å}$ b = 7.591 (7) ÅCell parameters from 51 reflections $\theta = 4.5 - 11.9^{\circ}$ c = 8.523 (8) Å $\alpha = 99.480 (14)^{\circ}$ $\mu = 0.13 \text{ mm}^{-1}$ T = 296 K $\beta = 108.748 (13)^{\circ}$ $\gamma = 99.240 \ (14)^{\circ}$ Plate, colourless $V = 443.3 (7) \text{ Å}^3$ $0.25\times0.19\times0.08~mm$

Data collection

Bruker SMART APEX CCD diffractometer

1535 independent reflections

Radiation source: sealed tube 1025 reflections with $I > 2\sigma(I)$

graphite $R_{\text{int}} = 0.025$

 ω and ϕ scans $\theta_{max} = 25.0^{\circ}, \, \theta_{min} = 2.6^{\circ}$

Absorption correction: multi-scan (SADABS; Bruker, 2008) $h = -8 \rightarrow 8$

 $T_{\text{min}} = 0.969$, $T_{\text{max}} = 0.990$ $k = -9 \rightarrow 8$ 2526 measured reflections $l = -10 \rightarrow 10$

Refinement

Refinement on F^2 Primary atom site location: structure-invariant direct methods

Least-squares matrix: full Secondary atom site location: difference Fourier map

 $R[F^2 > 2\sigma(F^2)] = 0.066$ Hydrogen site location: inferred from neighbouring

20(1)] 0.000

 $wR(F^2) = 0.190$ H-atom parameters constrained

S = 1.02 $w = 1/[\sigma^2(F_0^2) + (0.P)^2 + 0.1145P]$

where $P = (F_0^2 + 2F_c^2)/3$

1535 reflections $(\Delta/\sigma)_{max} = 0.001$

128 parameters $\Delta \rho_{max} = 0.24 \ e \ \text{Å}^{-3}$

0 restraints $\Delta \rho_{min} = -0.22 \text{ e Å}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F, with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F, and R- factors based on ALL data will be even larger.

H atoms were placed geometrically and allowed to refine as atoms riding on their bonding partners. The hydrogen was placed on the carboxylic acid based on the longer of the carboxylic acid carbon-oxygen bonds.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(\mathring{\mathbb{A}}^2)$

	x	y	Z	$U_{\rm iso}*/U_{\rm eq}$
F1	0.4739 (2)	1.3542 (2)	0.6014(2)	0.0629 (5)
O1	0.8299 (2)	1.4232 (4)	0.5841 (2)	0.0672 (7)
O2	0.8296 (2)	1.3413 (2)	0.3192 (2)	0.0716 (8)
H2A	0.9326	1.4171	0.3603	0.107*
O3	-0.1230 (4)	0.8633 (4)	0.2593 (2)	0.0798 (9)
O4	-0.0485 (2)	0.7256 (2)	0.0437 (2)	0.0582 (7)
C1	0.5568 (4)	1.2070 (4)	0.3734 (2)	0.0453 (7)
C2	0.4248 (4)	1.2212 (4)	0.4562 (2)	0.0464 (7)

C3	0.2443 (4)	1.1083 (4)	0.3967	(2)	0.0479 (8)	
НЗА	0.1607	1.1227	0.4553		0.057*	
C4	0.1871 (4)	0.9720 (4)	0.2478		0.0437 (7)	
C5	0.3166 (4)	0.9507 (4)	0.1625		0.0495 (8)	
H5A	0.2807	0.8584	0.0642		0.059*	
C6	0.4976 (5)	1.0670 (4)	0.2248		0.0527 (8)	
Н6А	0.5819	1.0523	0.1669		0.063*	
C7	0.7527 (4)	1.3333 (4)	0.4321		0.0503 (8)	
C8	-0.0112 (4)	0.8498 (4)	0.1877		0.0491 (8)	
C9	-0.2360 (5)	0.5972 (5)	-0.021		0.0692 (10)	
Н9А	-0.2482	0.5131	-0.123	` '	0.104*	
H9B	-0.3359	0.6639	-0.046		0.104*	
H9C	-0.2474	0.5303	0.0623	2	0.104*	
1170	0.2474	0.5505	0.0023		0.104	
		2				
Atomic displace	ment parameters	(\mathring{A}^2)				
	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
F1	0.0579 (11)	0.0628 (13)	0.0540 (10)	-0.0020 (9)	0.0211 (9)	-0.0111 (8)
O1	0.0526 (15)	0.0782 (17)	0.0532 (13)	-0.0072 (11)	` '	0.0038 (11)
O2	0.0590 (16)	0.0765 (18)	0.0690 (15)	-0.0133 (11)		0.0019 (11)
O3	0.0532 (16)	0.088 (2)	0.0828 (17)	-0.0110 (13)		-0.0154 (14)
O4	0.0485 (14)	0.0591 (14)	0.0525 (11)	-0.0015 (10		-0.0038 (10)
C1	0.0435 (17)	0.0453 (17)	0.0477 (15)	0.0102 (14)	0.0161 (13)	0.0123 (13)
C2	0.0485 (18)	0.0445 (17)	0.0414 (14)	0.0071 (13)	0.0153 (13)	0.0026 (11)
C3	0.0447 (18)	0.051 (2)	0.0470 (15)	0.0086 (14)	0.0202 (13)	0.0045 (13)
C4	0.0431 (17)	0.0430 (16)	0.0434 (15)	0.0074 (13)	0.0149 (13)	0.0091 (11)
C5	0.050 (2)	0.0467 (18)	0.0490 (16)	0.0076 (14)	0.0208 (14)	0.0017 (13)
C6	0.050 (2)	0.056 (2)	0.0535 (17)	0.0099 (15)	0.0243 (14)	0.0063 (14)
C7	0.049 (2)	0.0493 (18)	0.0510 (17)	0.0095 (14)	0.0168 (15)	0.0105 (14)
C8	0.0432 (18)	0.0511 (18)	0.0490 (16)	0.0079 (14)	0.0156 (14)	0.0055 (13)
C9	0.052 (2)	0.062 (2)	0.069 (2)	-0.0055 (17	* *	-0.0018 (17)
	()	()	,		,	
Coomotuio nava	mataus (Å °)					
Geometric para	meiers (A,)	1.251.(2)				22 (1)
F1—C2		1.364 (3)	C3—C			93 (4)
O1—C7		1.257 (3)	C3—H		0.93	
O2—C7		1.278 (4)	C4—C5		1.405 (4)	
O2—H2A		0.82	C4—C8		1.504 (4)	
O3—C8		1.197 (4)	C5—C			84 (4)
O4—C8		1.336 (4)	C5—H		0.93	
O4—C9		1.460 (4)	C6—H		0.93	
C1—C2		1.400 (4)	С9—Н		0.90	
C1—C6		1.405 (4)	C9—H		0.90	
C1—C7		1.504 (4)	С9—Н	9C	0.90	5
C2—C3		1.372 (4)				

C4—C5—H5A

C5—C6—C1

C5—C6—H6A

120.0

119.3

121.4 (3)

C7—O2—H2A

C8—O4—C9

C2—C1—C6

109.5

115.8 (2)

116.8 (3)

C2—C1—C7	124.1 (3)	C1—C6—H6A	119.3
C6—C1—C7	119.2 (2)	O1—C7—O2	124.2 (3)
F1—C2—C3	117.5 (2)	O1—C7—C1	119.9 (3)
F1—C2—C1	119.5 (3)	O2—C7—C1	115.9 (3)
C3—C2—C1	122.9 (3)	O3—C8—O4	124.0 (3)
C2—C3—C4	119.5 (3)	O3—C8—C4	124.0 (3)
C2—C3—H3A	120.2	O4—C8—C4	112.0(2)
C4—C3—H3A	120.2	O4—C9—H9A	109.5
C3—C4—C5	119.4 (3)	O4—C9—H9B	109.5
C3—C4—C8	117.9 (2)	H9A—C9—H9B	109.5
C5—C4—C8	122.7 (3)	O4—C9—H9C	109.5
C6—C5—C4	120.0 (3)	H9A—C9—H9C	109.5
C6—C5—H5A	120.0	H9B—C9—H9C	109.5

Hydrogen-bond geometry (Å, °)

D— H ··· A	<i>D</i> —H	$H\cdots A$	D··· A	$D\!\!-\!$
C9—H9A···F1 ⁱ	0.96	2.54	3.278 (5)	134 (1)
O2 ⁱⁱ —H2A ⁱⁱ ···O1	0.82	1.86	2.672 (4)	170 (1)
C3—H3A···O3 ⁱⁱⁱ	0.93	2.53	3.325 (4)	144 (1)

Symmetry codes: (i) x-1, y-1, z-1; (ii) -x+2, -y+3, -z+1; (iii) -x, -y+2, -z+1.

Fig. 1

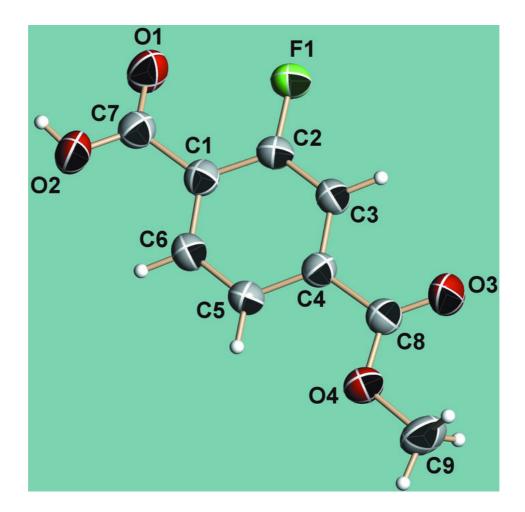


Fig. 2

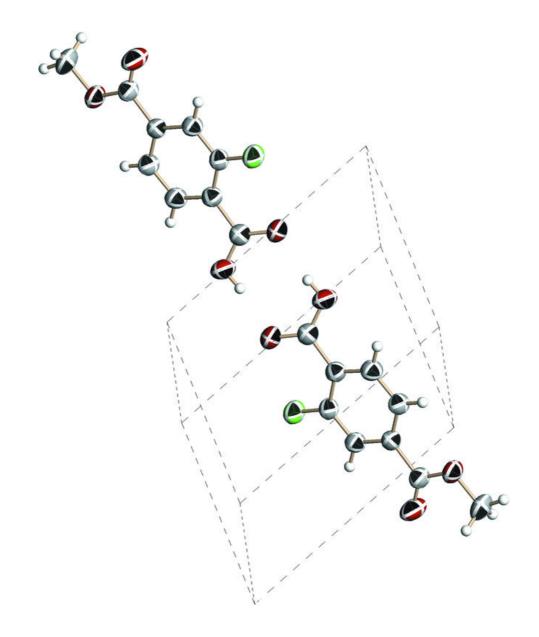


Fig. 3

