

Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: MU1071). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Hydrogen-Bonded Trimers of Methyl 4-Trifluoromethylpyrazole-3-carboxylate Solvated by Chloroform

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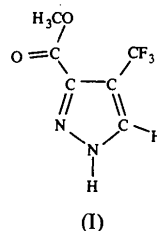
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Abstract

The title compound, C₆H₅F₃N₂O₂· $\frac{1}{6}$ CHCl₃, crystallizes from chloroform with three molecules of the ester and half a disordered chloroform molecule in the asymmetric unit. The three ester molecules form a hydrogen-bonded trimer *via* a cyclic arrangement of three N1—H···N2 hydrogen bonds. Corresponding bond lengths in all three molecules are similar.

Comment

The title pyrazole derivative (I) along with the 3- and 5-carbomethoxy isomers of 1-methyl-4-trifluoropyrazole are formed by the reaction of diazomethane and 4,4,4-trifluorobut-1-ynoic acid (Tajammal, 1988, 1991; Tajammal & Tipping, 1990). After chromatographic separation of the mixture of pyrazole derivatives, the title compound was recrystallized from chloroform, which supplies one solvate molecule for every six pyrazole molecules.



In the absence of 1-substitution, the N—H group at position 1 is available for hydrogen bonding and the novel trimerization seen in Fig. 1 occurs. The similar bond lengths in the pyrazole rings of the three molecules are in the range of lengths observed in 1-substituted pyrazoles (Beagley, Brown, Pritchard, Tajammal & Tipping, 1994). The N1···N2 distances for the three hydrogen bonds are N11···N32 = 2.902 (5), N21···N12 = 2.874 (6) and N31···N22 = 2.964 (5) Å. The third of these is significantly longer than the other two and is associated with a short N1 to carbonyl O atom distance,

$N31 \cdots O22 = 2.978(5) \text{ \AA}$, whereas the other two corresponding $N1 \cdots O2$ distances are greater than 3.1 \AA ; $O22$ belongs to a molecule in which the plane of the carboxylate group is rotated $11(1)^\circ$ from the pyrazole plane about the $C23\text{---}CO_2$ bond, whereas in the other two molecules the pyrazole rings and the carboxylate groups are approximately coplanar. Furthermore, the refined positions of the H atoms H11, H21 and H31 of the three $N1\text{---}H1 \cdots O2$ hydrogen bonds lead to a $H31 \cdots O22$ distance of $2.36(4) \text{ \AA}$, which is significantly shorter than the $H11 \cdots O32$ and $H21 \cdots O12$ distances of $2.68(4)$ and $2.77(4) \text{ \AA}$, respectively. However, all three $H1 \cdots N2$ distances are similar (about 2.0 \AA). The primary hydrogen bonding is thus between N1 and N2 in all three cases, but for $N31\text{---}H31 \cdots N22$ there is the additional weak participation of $O22$.

Each disordered chloroform solvate molecule lies across a twofold axis; Cl2 lies on the axis and Cl1, C1 and H1 are related to the equivalent atoms Cl1*, C1* and H1* by rotation about the axis. The three Cl-atom sites Cl1, Cl1* and Cl2 are fully occupied but the C—H component has an equal chance of being above or below the plane of the three Cl atoms.

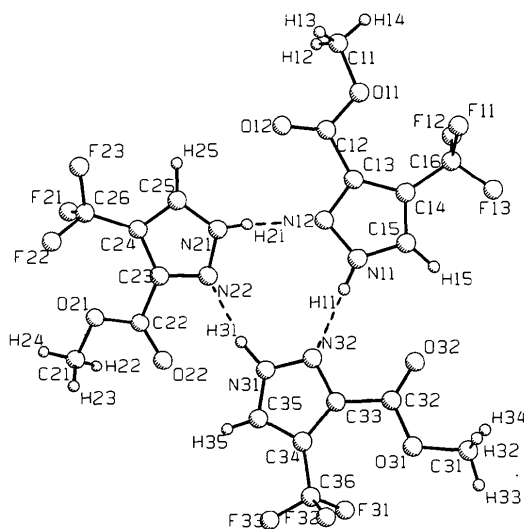


Fig. 1. The trimer, showing hydrogen bonding and atom labelling.

Experimental

Crystal data

$C_6H_5F_3N_2O_2 \cdot \frac{1}{6}CHCl_3$

$M_r = 214.05$

Monoclinic

$C2/c$

$a = 22.845(4) \text{ \AA}$

$b = 8.790(2) \text{ \AA}$

$c = 27.948(3) \text{ \AA}$

$\beta = 108.38(4)^\circ$

$V = 5326(4) \text{ \AA}^3$

Mo $K\alpha$ radiation

$\lambda = 0.71069 \text{ \AA}$

Cell parameters from 20 reflections

$\theta = 6.3\text{--}11.3^\circ$

$\mu = 0.30 \text{ mm}^{-1}$

$T = 296 \text{ K}$

Block

$0.4 \times 0.2 \times 0.2 \text{ mm}$

$Z = 24$

$D_x = 1.602 \text{ Mg m}^{-3}$

Data collection

Enraf-Nonius CAD-4 diffractometer

$\omega/2\theta$ scans

Absorption correction: none

7479 measured reflections

4490 independent reflections

2220 observed reflections

$[I > 3\sigma(I)]$

Colourless

$R_{int} = 0.022$

$\theta_{max} = 23^\circ$

$h = 0 \rightarrow 24$

$k = -9 \rightarrow 9$

$l = -27 \rightarrow 28$

3 standard reflections

frequency: 120 min

intensity variation:

insignificant

Refinement

Refinement on F

$R = 0.045$

$wR = 0.049$

$S = 2.09$

2220 reflections

403 parameters

Default weighting scheme in TEXSAN used

$(\Delta/\sigma)_{max} = 0.01$

$\Delta\rho_{max} = 0.23 \text{ e \AA}^{-3}$

$\Delta\rho_{min} = -0.33 \text{ e \AA}^{-3}$

Atomic scattering factors

from *International Tables*

for X-ray Crystallography

(1974, Vol. IV)

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (\AA^2)

$$B_{eq} = (8\pi^2/3) \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>B</i> _{eq}
Cl1	-0.06304 (9)	-0.4408 (3)	0.24668 (7)	11.7 (1)
Cl2	0	-0.7228 (3)	1/4	14.1 (2)
Cl1†	0.0019 (6)	-0.526 (2)	0.2711 (4)	8.3 (6)
F11	0.28084 (14)	0.1552 (4)	0.47050 (13)	9.1 (2)
F12	0.25734 (13)	-0.0379 (4)	0.50662 (14)	9.5 (2)
F13	0.24823 (13)	-0.0477 (5)	0.42867 (13)	10.5 (2)
O11	0.36067 (14)	0.1217 (4)	0.57535 (11)	5.7 (2)
O12	0.4617 (2)	0.0719 (4)	0.60372 (13)	6.3 (2)
N11	0.4260 (2)	-0.1743 (4)	0.4753 (2)	4.8 (2)
N12	0.44881 (15)	-0.1030 (4)	0.51948 (14)	4.5 (2)
C11	0.3649 (2)	0.2105 (5)	0.6197 (2)	7.0 (3)
C12	0.4122 (2)	0.0602 (5)	0.5726 (2)	4.5 (2)
C13	0.4006 (2)	-0.0284 (5)	0.5254 (2)	4.0 (2)
C14	0.3475 (2)	-0.0553 (5)	0.4843 (2)	4.4 (2)
C15	0.3660 (2)	-0.1487 (6)	0.4536 (2)	5.0 (2)
C16	0.2841 (2)	0.0035 (8)	0.4735 (2)	6.6 (3)
F21	0.7222 (2)	-0.2300 (4)	0.72911 (13)	10.1 (2)
F22	0.76526 (15)	-0.1268 (4)	0.68106 (13)	10.1 (2)
F23	0.7058 (2)	0.0011 (4)	0.71174 (14)	11.2 (2)
O21	0.73169 (13)	-0.4458 (4)	0.64879 (11)	5.7 (1)
O22	0.67120 (13)	-0.5013 (4)	0.57105 (12)	5.6 (2)
N21	0.5697 (2)	-0.1461 (5)	0.5906 (2)	4.9 (2)
N22	0.59417 (15)	-0.2716 (4)	0.57745 (13)	4.4 (2)
C21	0.7665 (2)	-0.5845 (6)	0.6489 (2)	7.1 (3)
C22	0.6846 (2)	-0.4209 (5)	0.6077 (2)	4.3 (2)
C23	0.6488 (2)	-0.2851 (5)	0.6127 (2)	4.1 (2)
C24	0.6584 (2)	-0.1673 (5)	0.6478 (2)	4.6 (2)
C25	0.6066 (2)	-0.0802 (6)	0.6320 (2)	5.0 (2)
C26	0.7125 (3)	-0.1313 (7)	0.6918 (2)	6.8 (3)
F31	0.52214 (15)	-0.7448 (3)	0.34987 (12)	7.6 (2)
F32	0.56786 (14)	-0.5597 (3)	0.32811 (10)	7.5 (2)
F33	0.61771 (14)	-0.7121 (4)	0.38575 (12)	8.7 (2)
O31	0.43935 (14)	-0.4882 (4)	0.32301 (12)	6.4 (2)
O32	0.40743 (14)	-0.3199 (4)	0.36929 (12)	6.4 (2)
N31	0.5655 (2)	-0.4315 (4)	0.47938 (15)	4.5 (2)
N32	0.5108 (2)	-0.3805 (4)	0.44977 (13)	4.4 (2)
C31	0.3843 (2)	-0.4627 (8)	0.2804 (2)	8.7 (3)
C32	0.4447 (2)	-0.4096 (6)	0.3646 (2)	4.8 (2)
C33	0.5026 (2)	-0.4474 (4)	0.4054 (2)	3.9 (2)

C34	0.5526 (2)	-0.5422 (4)	0.4072 (2)	4.1 (2)
C35	0.5918 (2)	-0.5271 (5)	0.4552 (2)	4.6 (2)
C36	0.5646 (2)	-0.6392 (6)	0.3679 (2)	5.4 (2)

† Site occupancy = 0.5.

Table 2. Selected geometric parameters (Å, °)

C11—C1	1.61 (1)	O22—C22	1.202 (5)
C11 ¹ —C1	1.79 (1)	N21—N22	1.340 (5)
C12—C1	1.83 (1)	N21—C25	1.330 (6)
F11—C16	1.337 (6)	N22—C23	1.329 (5)
F12—C16	1.312 (6)	C22—C23	1.479 (6)
F13—C16	1.339 (6)	C23—C24	1.395 (6)
O11—C11	1.443 (6)	C24—C25	1.361 (6)
O11—C12	1.320 (5)	C24—C26	1.477 (6)
O12—C12	1.194 (5)	F31—C36	1.323 (5)
N11—N12	1.336 (5)	F32—C36	1.336 (5)
N11—C15	1.332 (5)	F33—C36	1.322 (5)
N12—C13	1.336 (5)	O31—C31	1.451 (6)
C12—C13	1.483 (6)	O31—C32	1.324 (5)
C13—C14	1.401 (6)	O32—C32	1.198 (5)
C14—C15	1.348 (6)	N31—N32	1.339 (4)
C14—C16	1.477 (6)	N31—C35	1.335 (5)
F21—C26	1.319 (6)	N32—C33	1.332 (5)
F22—C26	1.331 (6)	C32—C33	1.485 (6)
F23—C26	1.320 (6)	C33—C34	1.402 (5)
O21—C21	1.455 (6)	C34—C35	1.364 (6)
O21—C22	1.321 (5)	C34—C36	1.481 (6)
C11 ¹ —C1—C11	112.7 (8)	C22—C23—C24	134.2 (4)
C11—C1—C12	112.3 (7)	C23—C24—C25	105.1 (4)
C11 ¹ —C1—C12	104.3 (6)	C23—C24—C26	130.5 (4)
C11—O11—C12	116.9 (4)	C25—C24—C26	124.3 (4)
N12—N11—C15	112.7 (4)	N21—C25—C24	106.6 (5)
N11—N12—C13	104.4 (3)	F21—C26—F22	105.1 (5)
O11—C12—O12	125.5 (4)	F21—C26—F23	105.0 (5)
O11—C12—C13	110.7 (4)	F21—C26—C24	113.9 (5)
O12—C12—C13	123.8 (4)	F22—C26—F23	107.7 (5)
N12—C13—C12	116.4 (4)	F22—C26—C24	113.4 (4)
N12—C13—C14	110.5 (4)	F23—C26—C24	111.2 (5)
C12—C13—C14	133.1 (4)	C31—O31—C32	117.2 (4)
C13—C14—C15	105.1 (4)	N32—N31—C35	112.2 (4)
C13—C14—C16	130.5 (4)	N31—N32—C33	105.0 (3)
C15—C14—C16	124.5 (4)	O31—C32—O32	124.5 (4)
N11—C15—C14	107.3 (5)	O31—C32—C33	111.7 (4)
F11—C16—F12	106.9 (5)	O32—C32—C33	123.7 (4)
F11—C16—F13	105.7 (5)	N32—C33—C32	116.5 (4)
F11—C16—C14	113.2 (5)	N32—C33—C34	110.8 (4)
F12—C16—F13	106.4 (5)	C32—C33—C34	132.8 (4)
F12—C16—C14	113.8 (5)	C33—C34—C35	106.4 (4)
F13—C16—C14	110.3 (4)	C33—C34—C36	130.9 (4)
C21—O21—C22	115.8 (4)	C35—C34—C36	124.5 (4)
N22—N21—C25	113.1 (4)	N31—C35—C34	107.4 (4)
N21—N22—C23	104.2 (4)	F31—C36—F32	106.0 (4)
O21—C22—O22	124.2 (4)	F31—C36—F33	106.2 (4)
O21—C22—C23	112.0 (4)	F31—C36—C34	113.7 (4)
O22—C22—C23	123.8 (4)	F32—C36—F33	106.4 (4)
N22—C23—C22	114.8 (4)	F32—C36—C34	112.8 (4)
N22—C23—C24	111.0 (4)	F33—C36—C34	111.3 (4)

Symmetry code: (i) $-x, y, \frac{1}{2} - z$.

The data were collected in terms of a triclinic cell and later transformed and merged to give Miller indices of the reported equivalent centred monoclinic cell of twice the volume. For each of the three substituted pyrazole molecules, the positions of the H atoms H1 and H5 attached to the rings were refined isotropically. Methyl H atoms were placed in calculated positions. Data collection: *CAD-4 Software* (Enraf-Nonius, 1989). Cell refinement: *CAD-4 Software*. Data reduction: *SHELX76* (Sheldrick, 1976). Program(s) used to solve structure: *MITHRIL* (Gilmore, 1984); *SHELXS86* (Sheldrick, 1985). Program(s) used to refine structure: *TEXSAN* (Molecular Structure Corporation, 1985). Molecular graphics: *PLUTO* (Moth-

erwell & Clegg, 1978). Software used to prepare material for publication: *CIF* (Hall, Allen & Brown, 1991).

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: HA1076). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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p-Hydroxyephedrinium Dihydrogenphosphate

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Abstract

The structural investigation of the title compound, [1-hydroxy-1-(4-hydroxyphenyl)-2-propyl]-methylammonium dihydrogenphosphate, C₁₀H₁₆NO₂⁺·H₂PO₄⁻, has been carried out by single-crystal X-ray diffraction. The ethylamine side chain