



Figure 3
A projection of the difference Fourier map for (I) in the plane of the main F atoms, showing the disordered charge (when disorder is not accounted for).

The ionization states of the glycine and trifluoroacetic acid molecules in (I) were determined from the objective localization of the H atoms bonded to the carboxylic acid groups, but could easily be inferred from the bond distances within these groups. The glycine molecule exists in the cationic form, with a mono-positively charged amino group and a neutral carboxylic acid group, in agreement with the large asymmetry between the C—O bond lengths of this functional group. The trifluoroacetate molecules are in the ionized state, as expected from the strength of the acid and the required charge neutrality of the salt.

The glycine carboxyl skeleton, which includes atoms O1, O2, C1 and C2, is planar to within 0.0010 (4) Å. The N atom is slightly displaced out of this plane, by -0.072 (3) Å, corresponding to a small rotation around the C1—C2 single bond. The relevant torsion angles are O1—C1—C2—N of 3.3 (2)° and O2—C1—C2—N of -177.09 (13)°. These can be compared with the corresponding values in pure γ -glycine [167.1 (1) and -15.4 (1)°, respectively; Kvick *et al.*, 1980], which is more distorted from planarity.

The trifluoroacetate anion in (I) has a staggered conformation, as indicated by the O—C4—C5—F torsion angles. The geometry of the CF₃ group is similar to that found in other structures (Nahrngbauer *et al.*, 1979), with an average C—F bond length and F—C—F angle of 1.327 (7) Å and 106.8 (2)°, respectively. The average F—C—C angle is 112 (2)°. The carboxylate group of the anion is planar to within 0.0089 (4) Å; the C4—C5 bond length of 1.541 (2) Å is longer than the average value for a Csp³—Csp² bond (Allen *et al.*, 1987), but is within the normal range of values found in trifluoroacetic acid and trifluoroacetate compounds (Lundgren, 1978). The CF₃ group is disordered over two very close positions, the most probable occurring with roughly 0.75 probability. Low-temperature measurements seem to support the possible dynamic nature of this disorder, as no refinement of two different positions for each F atom is then necessary.

The glycine cations in (I) interact directly, *via* hydrogen bonds, with two neighbouring trifluoroacetate anions, forming cation–anion–cation–anion rings lying in the *bc* plane. The glycine cation acts as a donor in four hydrogen bonds; the N atom is engaged in three hydrogen bonds as a donor, and the O atom of the carboxylic acid group is also the donor of the strongest hydrogen bond found in this structure. Both O atoms of the carboxylate group of the anion act as acceptors of hydrogen bonds. The above-mentioned ring is formed *via* two very strong O—H···O and two weaker N—H···O hydrogen bonds. The N—H···O distances and angles are in the ranges 2.856 (2)– 2.984 (2) Å and 159.2 (18)– 175.5 (17)°, respectively, which allows a classification of these hydrogen bonds as relatively weak (Desiraju & Steiner, 1999). The strongest hydrogen bond in (I) involves the carboxylic acid O atom of the cation as a donor and a carboxylate O atom of the anion as an acceptor [O—H···O = 2.597 (2) Å and O—H—O = 175 (2)°].

Experimental

Colourless prismatic crystals of (I) were obtained by evaporation of the solvent from a solution of glycine in trifluoroacetic acid. Good quality single crystals recrystallized after a few hours at room temperature. A small single crystal was cast and checked by photographic methods prior to data collection.

Crystal data

C ₂ H ₆ NO ₂ ⁺ ·C ₂ F ₃ O ₂ [−]	$D_x = 1.798$ Mg m ^{−3}
$M_r = 189.10$	Mo K α radiation
Monoclinic, $P2_1/c$	Cell parameters from 25 reflections
$a = 4.9598$ (6) Å	$\theta = 9.6$ – 15.4 °
$b = 12.239$ (2) Å	$\mu = 0.20$ mm ^{−1}
$c = 12.015$ (8) Å	$T = 293$ (2) K
$\beta = 106.67$ (7)°	Prismatic, colourless
$V = 698.7$ (5) Å ³	$0.47 \times 0.40 \times 0.15$ mm
$Z = 4$	

Table 1

Selected geometric parameters (Å, °).

O1—C1	1.3127 (17)	F2—C4	1.313 (6)
O2—C1	1.1997 (17)	F3—C4	1.323 (7)
N—C2	1.4736 (18)	O4—C3	1.2446 (17)
C1—C2	1.5084 (18)	O3—C3	1.2317 (17)
F1—C4	1.361 (7)	C3—C4	1.5430 (19)
O2—C1—O1	126.11 (12)	O4—C3—C4	114.21 (11)
O2—C1—C2	122.61 (12)	F2—C4—C3	114.0 (3)
O1—C1—C2	111.28 (11)	F3—C4—C3	109.0 (4)
O3—C3—O4	129.44 (12)	F1—C4—C3	112.2 (3)
O3—C3—C4	116.28 (12)		

Table 2

Hydrogen-bonding geometry (Å, °).

<i>D</i> —H··· <i>A</i>	<i>D</i> —H	H··· <i>A</i>	<i>D</i> ··· <i>A</i>	<i>D</i> —H··· <i>A</i>
O1—H1···O4 ⁱ	0.86 (2)	1.74 (3)	2.597 (2)	175 (2)
N—H5···O4 ⁱⁱ	0.93 (2)	1.93 (2)	2.856 (2)	175.5 (17)
N—H4···O3 ⁱⁱⁱ	0.91 (2)	2.04 (2)	2.941 (2)	173.0 (16)
N—H6···O3 ^{iv}	0.93 (2)	2.09 (2)	2.984 (2)	159.2 (18)

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

Data collection

Enraf–Nonius CAD-4 diffractometer	$\theta_{\max} = 27.5^\circ$
Profile data from $\omega/2\theta$ scans	$h = -6 \rightarrow 0$
1772 measured reflections	$k = -15 \rightarrow 0$
1599 independent reflections	$l = -14 \rightarrow 15$
1285 reflections with $I > 2\sigma(I)$	25 standard reflections
$R_{\text{int}} = 0.021$	frequency: 180 min
	intensity decay: 7%

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0538P)^2 + 0.1642P]$
$R[F^2 > 2\sigma(F^2)] = 0.030$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.092$	$(\Delta/\sigma)_{\max} < 0.001$
$S = 1.02$	$\Delta\rho_{\max} = 0.32 \text{ e } \text{\AA}^{-3}$
1599 reflections	$\Delta\rho_{\min} = -0.19 \text{ e } \text{\AA}^{-3}$
154 parameters	Extinction correction: <i>SHELXL97</i>
H atoms treated by a mixture of independent and constrained refinement	(Sheldrick, 1997)
	Extinction coefficient: 0.197 (10)

All H atoms could be located in a difference Fourier map; those attached to atom C2 were placed in idealized positions and refined as riding, while the remaining H atoms were refined isotropically [$\text{O}-\text{H} = 0.86$ (2) \AA and $\text{N}-\text{H} = 0.91$ (2)– 0.93 (2) \AA]. Examination of the structure with *PLATON* (Spek, 2001) showed that there were no solvent-accessible voids in the crystal lattice.

Data collection: *CAD-4 Software* (Enraf–Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *PLATON* (Spek, 2001); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976); software used to prepare material for publication: *SHELXL97*.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: SK1592). Services for accessing these data are described at the back of the journal.

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