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Key indicators

Single-crystal X-ray study
 $T = 297\text{ K}$
Mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$
 R factor = 0.040
 wR factor = 0.143
Data-to-parameter ratio = 12.2For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.Anhydrous guanidinium hydrogen fumarate:
a two-dimensional hydrogen-bonded network
structureThe title compound, $\text{CH}_6\text{N}_3^+\cdot\text{C}_4\text{H}_3\text{O}_4^-$, forms a two-dimensional hydrogen-bonded network structure. The guanidinium cations lie across crystallographic mirror planes while the hydrogen fumarate anions have inversion symmetry with the acid proton 50% disordered over the two carboxylate groups.Received 7 December 2006
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Comment

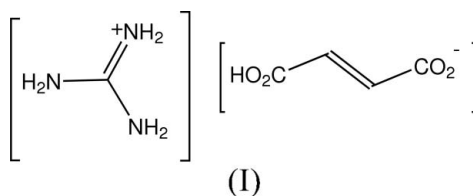
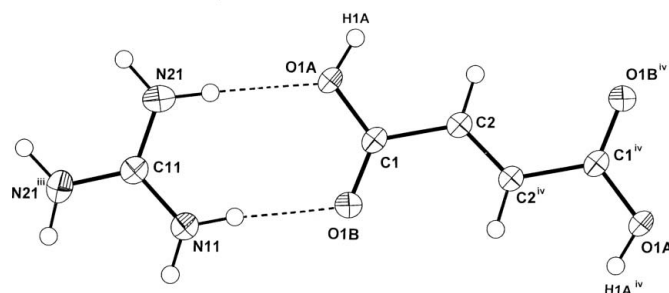
The known structures of guanidinium salts with diprotic aliphatic acids are not common and many are acid salts (Adams, 1978; Andrews *et al.*, 1979; Leban & Rupnik, 1992; Zyss *et al.*, 1993; Krumbe *et al.*, 1989; Krumbe & Haussuehl, 1987; Golic *et al.*, 1985; Froehlich *et al.*, 1985).Since the incorporation of water molecules of hydration in guanidinium salts is dependent on the choice of solvent, we carried out the synthesis of (I) in 50% propan-2-ol–water to obtain the anhydrous 1:1 salt reported here. The centrosymmetric $P2_1/m$ model (rather than the $P2_1$ model which was tried but gave poor refinement and low precision) places the guanidinium cation across a crystallographic mirror plane and the hydrogen fumarate anion across an inversion centre. This requires the single acid proton to be 50% disordered over the

Figure 1

Molecular structure and atom-naming scheme for the guanidinium cation and the hydrogen fumarate anion in (I). Displacement ellipsoids are drawn at the 50% probability level. The guanidinium atoms N21 and N21ⁱⁱⁱ are related by mirror symmetry while the fumarate atoms designated (iv) are inversion-related [symmetry codes: (iii) $x, -y + \frac{1}{2}, z$; (iv) $-x, -y, -z + 1$]. The fumarate acid protons H1A and H1A^{iv} are 50% disordered over the two carboxylate groups. Dashed lines indicate hydrogen bonds.

two *trans*-related carboxylate groups (Fig. 1). All available proton donors and acceptors are involved in hydrogen-bonding interactions (Table 1), giving a two-dimensional layered network structure (Fig. 2). For the guanidinium cation, there are six cyclic associations [two $R_2^2(8)$, two $R_4^4(10)$ and two $R_5^5(11)$]. The two mirror-related 11-membered rings are closed by single *anti*-related fumaric acid H atoms which also give intermolecular *anti*-associations with fumarate-O acceptors in strong $O1A \cdots O1A^i$ hydrogen bonds (Table 1).

Experimental

The title compound was synthesized by heating together fumaric acid (1 mmol) and guanidinium carbonate (0.5 mmol) in 50 ml of 50% propan-2-ol-water under reflux for 10 min. After concentration to ca 30 ml, partial room-temperature evaporation of the hot-filtered solution gave colourless crystal plates (m.p. 495–497 K).

Crystal data

| | |
|----------------------------------|---|
| $CH_6N_3^+ \cdot C_4H_3O_4^-$ | $Z = 2$ |
| $M_r = 175.15$ | $D_x = 1.525 \text{ Mg m}^{-3}$ |
| Monoclinic, $P2_1/m$ | Mo $K\alpha$ radiation |
| $a = 5.4995$ (12) Å | $\mu = 0.13 \text{ mm}^{-1}$ |
| $b = 18.842$ (5) Å | $T = 297$ (2) K |
| $c = 3.6943$ (11) Å | Plate, colourless |
| $\beta = 94.83$ (2)° | $0.45 \times 0.32 \times 0.12 \text{ mm}$ |
| $V = 381.45$ (17) Å ³ | |

Data collection

| | |
|---------------------------------------|------------------------------------|
| Rigaku AFC-7R diffractometer | $R_{\text{int}} = 0.025$ |
| ω - 2θ scans | $\theta_{\text{max}} = 27.5^\circ$ |
| Absorption correction: none | 3 standard reflections |
| 1088 measured reflections | frequency: 150 min |
| 905 independent reflections | intensity decay: 2.4% |
| 637 reflections with $I > 2\sigma(I)$ | |

Refinement

| | |
|--|--|
| Refinement on F^2 | $w = 1/[\sigma^2(F_o^2) + (0.1P)^2 + 0.0662P]$ |
| $R[F^2 > 2\sigma(F^2)] = 0.040$ | where $P = (F_o^2 + 2F_c^2)/3$ |
| $wR(F^2) = 0.143$ | $(\Delta/\sigma)_{\text{max}} < 0.001$ |
| $S = 0.94$ | $\Delta\rho_{\text{max}} = 0.17 \text{ e \AA}^{-3}$ |
| 905 reflections | $\Delta\rho_{\text{min}} = -0.26 \text{ e \AA}^{-3}$ |
| 74 parameters | |
| H atoms treated by a mixture of independent and constrained refinement | |

Table 1

Hydrogen-bond geometry (Å, °).

| $D-H \cdots A$ | $D-H$ | $H \cdots A$ | $D \cdots A$ | $D-H \cdots A$ |
|----------------------------|----------|--------------|--------------|----------------|
| $O1A-H1A \cdots O1A^i$ | 0.85 (4) | 1.63 (4) | 2.4658 (18) | 166 (5) |
| $N11-H11 \cdots O1B$ | 0.88 (3) | 1.99 (3) | 2.873 (2) | 174 (2) |
| $N21-H21A \cdots O1B^{ii}$ | 0.92 (3) | 2.19 (3) | 3.006 (2) | 148 (3) |
| $N21-H21B \cdots O1A$ | 0.89 (3) | 2.10 (3) | 2.991 (2) | 175 (3) |

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

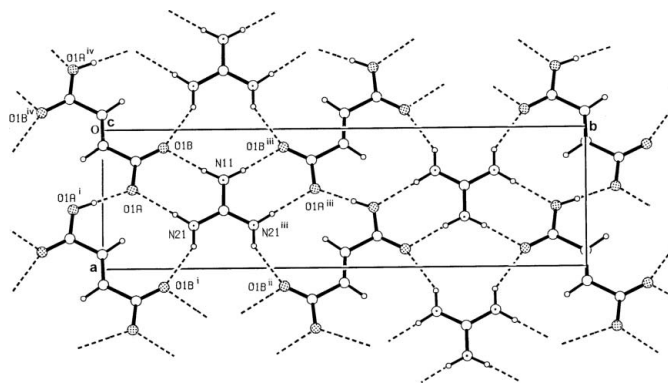


Figure 2

The hydrogen-bonded sheet structure of (I) viewed down the c axial direction, showing hydrogen-bonding associations as dashed lines. For symmetry codes, see Fig. 1 caption and Table 1.

The guanidinium H atoms and the fumarate acid H atom were located by difference methods and their positional and isotropic displacement parameters were refined. The fumarate ethylenic H atom was included in the refinement in a calculated position ($C-H = 0.94$ Å) using a riding-model approximation, with $U_{\text{iso}}(H) = U_{\text{eq}}(C)$.

Data collection: *MSC/AFC Diffractometer Control Software* (Molecular Structure Corporation, 1999); cell refinement: *MSC/AFC Diffractometer Control Software*; data reduction: *TEXSAN* for Windows (Molecular Structure Corporation, 1999); program(s) used to solve structure: *SIR92* (Altomare *et al.*, 1994); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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References

- Adams, J. M. (1978). *Acta Cryst.* **B34**, 1218–1220.
 Altomare, A., Casciarano, G., Giacovazzo, C., Guagliardi, A., Burla, M. C., Polidori, G. & Camalli, M. (1994). *J. Appl. Cryst.* **27**, 435.
 Andrews, L. C., Deroski, B. R. & Ricci, J. S. (1979). *J. Cryst. Mol. Struct.* **9**, 163–171.
 Froehlich, R., Mattern, G. & Krumbe, W. (1985). *Z. Kristallogr.* **172**, 315–317.
 Golic, L., Leban, I., Detoni, S., Orel, B. & Hadzi, D. (1985). *J. Crystallogr. Spectrosc. Res.* **15**, 215–228.
 Krumbe, W. & Haussuehl, S. (1987). *Z. Kristallogr.* **179**, 267–269.
 Krumbe, W., Haussuehl, S. & Froehlich, R. (1989). *Z. Kristallogr.* **187**, 309–318.
 Leban, I. & Rupnik, A. (1992). *Acta Cryst.* **C48**, 821–824.
 Molecular Structure Corporation (1999). *MSC/AFC Diffractometer Control Software and TEXSAN for Windows* (Version 1.06). MSC, The Woodlands, Texas, USA.
 Sheldrick, G. M. (1997). *SHELXL97*. University of Göttingen, Germany.
 Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.
 Zyss, J., Pecaut, J., Levy, J. P. & Masse, R. (1993). *Acta Cryst.* **B49**, 334–342.