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Key indicators

Single-crystal X-ray study
 $T = 296$ K
Mean $\sigma(\text{C}-\text{C}) = 0.005$ Å
 R factor = 0.066
 wR factor = 0.221
Data-to-parameter ratio = 14.5For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Triethanolammonium 2-formylbenzoate

The title compound, $\text{C}_6\text{H}_{16}\text{NO}_3^+ \cdot \text{C}_8\text{H}_5\text{O}_3^-$, contains edge-fused $R_2^2(12)R_2^2(10)R_2^2(12)$ motifs involving triethanolammonium and 2-carboxylbenzoate ions. These motifs are connected by $C(9)$ chains with $R_2^2(12)$ rings.

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Comment

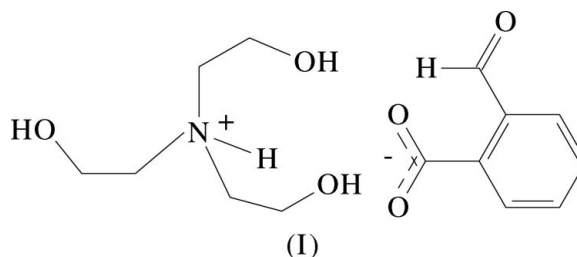
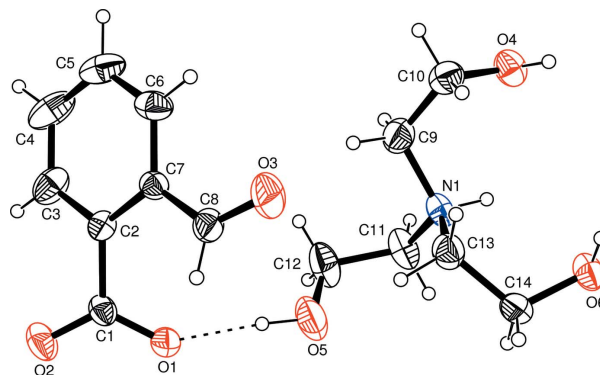
The title compound, (I), was obtained by the reaction of triethanolamine and 2-formylbenzoic acid. We have been interested in supramolecularly hydrogen-bonded systems formed by organic amines and carboxylic acids (Odabaşoğlu, Büyükgüngör & Lönnecke, 2003; Odabaşoğlu, Büyükgüngör *et al.*, 2003; Büyükgüngör *et al.*, 2004; Büyükgüngör & Odabaşoğlu, 2002, 2003, 2006; Odabaşoğlu & Büyükgüngör, 2006*a,b*). The present work is part of a structural study of compounds of organic ammonium systems with hydrogen-bond donors and we report here the structure of triethanolammonium 2-formylbenzoate, (I) (Fig. 1).In compound (I), the triethanolammonium ions are linked to the 2-formylbenzoate ions through $\text{N}-\text{H} \cdots \text{O}$ and $\text{O}-\text{H} \cdots \text{O}$ hydrogen bonds (Table 1) and generate centrosymmetric edge-fused $[R_2^2(12)R_2^2(10)R_2^2(12)]$ motifs (Fig. 2) (Etter,

Figure 1

The molecular structure of (I), with the atomic numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. The dashed line indicates a hydrogen bond.

1990). These motifs are connected by $C(9)$ chains with $R_2^2(12)$ rings (Fig. 3). Bond lengths and angles for the 2-formylbenzoate ion are similar to those reported in our previous work (Büyükgüngör & Odabaşoğlu, 2006).

Experimental

The title compound was prepared by mixing triethanolamine (1.49 g, 0.01 mol) and 2-formylbenzoic acid (1.50 g, 0.01 mol) in water (50 ml) at room temperature. Crystals of (I) were obtained by slow evaporation of the solvent (m.p. 354–357 K).

Crystal data

$C_6H_{16}NO_3^+ \cdot C_8H_5O_3^-$	$Z = 4$
$M_r = 299.32$	$D_x = 1.328 \text{ Mg m}^{-3}$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 11.5316 (8) \text{ \AA}$	$\mu = 0.10 \text{ mm}^{-1}$
$b = 7.5990 (4) \text{ \AA}$	$T = 296 \text{ K}$
$c = 19.2535 (14) \text{ \AA}$	Prism, colourless
$\beta = 117.470 (5)^\circ$	$0.76 \times 0.61 \times 0.38 \text{ mm}$
$V = 1496.93 (17) \text{ \AA}^3$	

Data collection

Stoe IPDS II diffractometer	13959 measured reflections
ω scans	2947 independent reflections
Absorption correction: integration (<i>X-RED32</i> ; Stoe & Cie, 2002)	2283 reflections with $I > 2\sigma(I)$
$T_{\min} = 0.752$, $T_{\max} = 0.949$	$R_{\text{int}} = 0.038$
	$\theta_{\text{max}} = 26.0^\circ$

Refinement

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.128P)^2 + 0.5183P]$
$R[F^2 > 2\sigma(F^2)] = 0.066$	where $P = (F_o^2 + 2F_c^2)/3$
$wR(F^2) = 0.221$	$(\Delta/\sigma)_{\text{max}} < 0.001$
$S = 1.10$	$\Delta\rho_{\text{max}} = 0.91 \text{ e \AA}^{-3}$
2947 reflections	$\Delta\rho_{\text{min}} = -0.33 \text{ e \AA}^{-3}$
203 parameters	Extinction correction: <i>SHELXL97</i>
H atoms treated by a mixture of independent and constrained refinement	(Sheldrick, 1997)
	Extinction coefficient: 0.008 (3)

Table 1

Hydrogen-bond geometry (\AA , $^\circ$).

$D-H \cdots A$	$D-H$	$H \cdots A$	$D \cdots A$	$D-H \cdots A$
$N1-H1 \cdots O6^i$	0.88 (4)	2.06 (3)	2.867 (3)	153 (3)
$O4-H4A \cdots O1^{ii}$	0.82 (4)	1.88 (4)	2.706 (3)	178 (3)
$O5-H5A \cdots O1$	0.92 (5)	1.79 (5)	2.709 (3)	175 (5)
$O6-H6A \cdots O2^{ii}$	0.84 (5)	1.82 (5)	2.645 (3)	168 (5)

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

All C-bound H atoms were refined using the riding-model approximation, with $C-H = 0.93 \text{ \AA}$ for aromatic and aldehydic H, and $C-H = 0.97 \text{ \AA}$ for methylene $C-H$ [$U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{parent atom})$]. The amino and hydroxy H atoms were located in a difference Fourier map and refined freely with an isotropic displacement parameter.

Data collection: *X-AREA* (Stoe & Cie, 2002); cell refinement: *X-AREA*; data reduction: *X-RED32* (Stoe & Cie, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1990); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP-3 for Windows* (Farrugia, 1997); software used to prepare material for publication: *WinGX* (Farrugia, 1999).

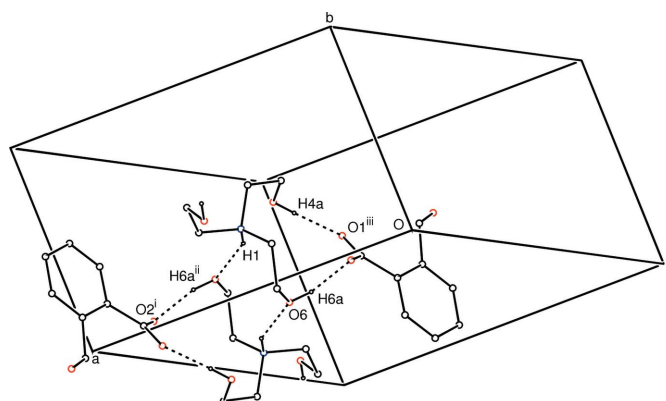


Figure 2

Part of the crystal structure of (I), showing the formation of a hydrogen-bonded $[R_2^2(12)R_2^2(10)R_2^2(12)]$ motif. H atoms not involved in hydrogen bonds have been omitted for clarity. [Symmetry codes: (i) $1 - x, y - \frac{1}{2}, z - \frac{1}{2}$; (ii) $1 - x, -y, -z$; (iii) $x - \frac{1}{2}, -y, z - \frac{1}{2}$].

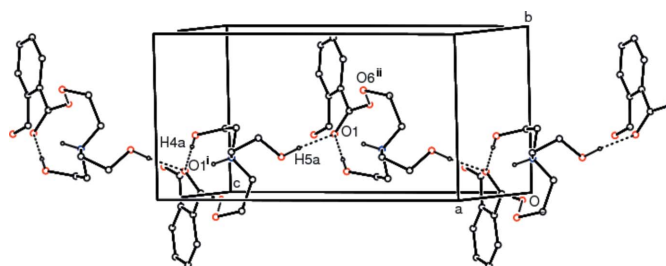


Figure 3

Part of the crystal structure of (I), showing the formation of a $C(9)$ chain with $R_2^2(12)$ ring. The $N-H \cdots O$ and $C-H \cdots O$ interactions are represented as dashed lines. H atoms not involved in hydrogen bonds have been omitted for clarity. [Symmetry codes: (i) $x, -y + \frac{1}{2}, z + \frac{1}{2}$; (ii) $x, -y + \frac{1}{2}, z - \frac{1}{2}$].

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