

**Bis(carboxymethyl)ammonium
4-toluenesulfonate****Kong Mun Lo and Seik Weng Ng***Department of Chemistry, University of Malaya, 50603 Kuala Lumpur, Malaysia
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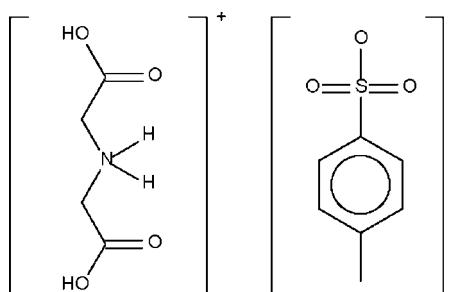
Received 8 August 2008; accepted 14 August 2008

Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C-C}) = 0.003 \text{ \AA}$; R factor = 0.038; wR factor = 0.119; data-to-parameter ratio = 15.4.

The iminodiacetic acid component of the title salt, $\text{C}_4\text{H}_8\text{NO}_4^+\cdot\text{C}_7\text{H}_7\text{SO}_3^-$, is protonated at the N atom. The cation uses the ammonium group to form hydrogen bonds to the O atoms of two adjacent sulfonate groups. In addition, the carboxylic acid portions of the cation form hydrogen bonds to the sulfonate groups. The hydrogen-bonding interactions give rise to a layer structure.

Related literature

For the crystal structures of iminodiacetic acid hydrohalides, see: Oskarsson (1973, 1974*a,b*, 1976).

**Experimental***Crystal data*
 $\text{C}_4\text{H}_8\text{NO}_4^+\cdot\text{C}_7\text{H}_7\text{O}_3\text{S}^-$
 $M_r = 305.30$
Orthorhombic, $Pbca$
 $a = 9.9291 (2) \text{ \AA}$
 $b = 10.3636 (2) \text{ \AA}$
 $c = 25.8862 (5) \text{ \AA}$
 $V = 2663.72 (9) \text{ \AA}^3$
 $Z = 8$
Mo $K\alpha$ radiation
 $\mu = 0.28 \text{ mm}^{-1}$ $T = 100 (2) \text{ K}$
 $0.27 \times 0.27 \times 0.27 \text{ mm}$ *Data collection*Bruker SMART APEX
diffractometer
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.929$, $T_{\max} = 0.929$ 20842 measured reflections
3059 independent reflections
2560 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$ *Refinement*
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.119$
 $S = 1.15$
3059 reflections
198 parameters
4 restraints

H atoms treated by a mixture of
independent and constrained
refinement
 $\Delta\rho_{\max} = 0.42 \text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.50 \text{ e \AA}^{-3}$
Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1—H1N1 \cdots O2 ⁱ	0.88 (1)	2.02 (1)	2.885 (2)	167 (2)
N1—H1N2 \cdots O3 ⁱⁱ	0.88 (1)	2.06 (2)	2.792 (2)	140 (2)
O5—H5O \cdots O1	0.84 (1)	1.79 (1)	2.607 (2)	164 (3)
O7—H7O \cdots O2 ⁱⁱⁱ	0.84 (1)	1.85 (1)	2.659 (2)	160 (3)

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *X-SEED* (Barbour, 2001); software used to prepare material for publication: *publCIF* (Westrip, 2008).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BH2187).

References

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supporting information

Acta Cryst. (2008). E64, o1798 [doi:10.1107/S1600536808026214]

Bis(carboxymethyl)ammonium 4-toluenesulfonate

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S1. Experimental

Iminodiacetic acid (0.55 g, 4 mmol) and *p*-toluenesulfonic acid (0.65 g, 4 mmol) were heated in toluene (100 ml) for 1 h. Crystals were isolated from the cool solution after several days.

S2. Refinement

Carbon-bound H-atoms were placed in calculated positions (C—H 0.95 to 0.99 Å) and were included in the refinement in the riding model approximation, with $U_{\text{iso}}(\text{H})$ set to 1.2 to 1.5 U_{eq} (carrier C). The acid and ammonium H atoms were refined with distance restraints of O—H = 0.84 (1) and N—H = 0.88 (1) Å; their isotropic displacement parameters were freely refined.

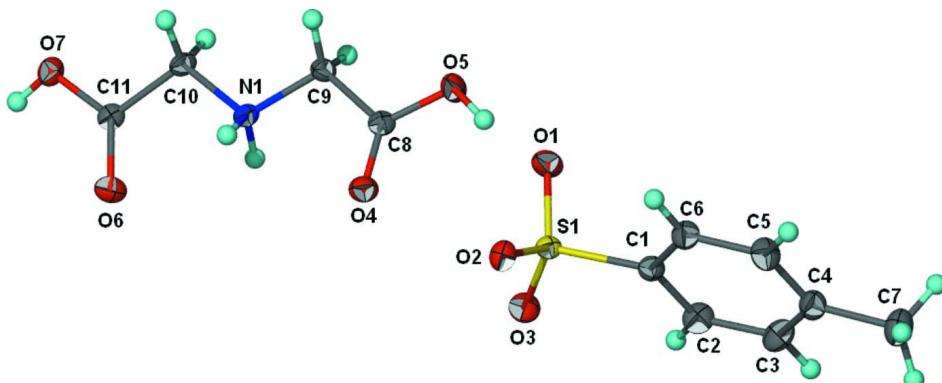
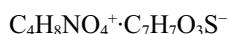


Figure 1

Thermal ellipsoid plot (Barbour, 2001) of $(\text{C}_4\text{H}_8\text{NO}_4)^+(\text{C}_7\text{H}_7\text{O}_3\text{S})^-$ at the 70% probability level. Hydrogen atoms are drawn as spheres of arbitrary radius.

Bis(carboxymethyl)ammonium 4-toluenesulfonate

Crystal data



$M_r = 305.30$

Orthorhombic, *Pbca*

Hall symbol: -P 2ac 2ab

$a = 9.9291 (2)$ Å

$b = 10.3636 (2)$ Å

$c = 25.8862 (5)$ Å

$V = 2663.72 (9)$ Å³

$Z = 8$

$F(000) = 1280$

$D_x = 1.523 \text{ Mg m}^{-3}$

Mo $\text{K}\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 4293 reflections

$\theta = 2.9\text{--}27.2^\circ$

$\mu = 0.28 \text{ mm}^{-1}$

$T = 100$ K

Triangular block, colorless

$0.27 \times 0.27 \times 0.27$ mm

Data collection

Bruker SMART APEX
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
 ω scans
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.930$, $T_{\max} = 0.930$

20842 measured reflections
3059 independent reflections
2560 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$
 $\theta_{\max} = 27.5^\circ$, $\theta_{\min} = 1.6^\circ$
 $h = -12 \rightarrow 7$
 $k = -13 \rightarrow 13$
 $l = -33 \rightarrow 33$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.038$
 $wR(F^2) = 0.119$
 $S = 1.15$
3059 reflections
198 parameters
4 restraints
0 constraints
Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
H atoms treated by a mixture of independent
and constrained refinement
 $w = 1/[\sigma^2(F_o^2) + (0.0644P)^2 + 0.8206P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} = 0.001$
 $\Delta\rho_{\max} = 0.42 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.50 \text{ e } \text{\AA}^{-3}$

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
S1	0.49041 (5)	0.84948 (4)	0.358654 (17)	0.01248 (14)
O1	0.55419 (15)	0.74172 (13)	0.38508 (5)	0.0195 (3)
O2	0.56209 (14)	0.88382 (13)	0.31102 (5)	0.0163 (3)
O3	0.34727 (14)	0.83294 (13)	0.35029 (5)	0.0190 (3)
O4	0.40223 (14)	0.52637 (13)	0.30853 (5)	0.0171 (3)
O5	0.51261 (16)	0.49339 (13)	0.38301 (5)	0.0199 (3)
H5O	0.510 (3)	0.5747 (10)	0.3841 (10)	0.034 (7)*
O6	0.20627 (14)	0.17528 (13)	0.21695 (5)	0.0176 (3)
O7	0.29656 (14)	-0.02406 (13)	0.22217 (5)	0.0163 (3)
H7O	0.229 (2)	-0.047 (3)	0.2048 (10)	0.051 (9)*
N1	0.37466 (17)	0.27383 (15)	0.28845 (6)	0.0129 (3)
H1N1	0.402 (3)	0.316 (2)	0.2610 (7)	0.031 (7)*
H1N2	0.2899 (11)	0.296 (2)	0.2924 (8)	0.014 (5)*
C1	0.50864 (19)	0.98391 (18)	0.40013 (7)	0.0141 (4)
C2	0.3995 (2)	1.02980 (19)	0.42797 (8)	0.0187 (4)
H2	0.3146	0.9879	0.4256	0.022*
C3	0.4152 (2)	1.1376 (2)	0.45941 (8)	0.0212 (4)
H3	0.3404	1.1681	0.4788	0.025*
C4	0.5374 (2)	1.20164 (19)	0.46306 (7)	0.0183 (4)
C5	0.6471 (2)	1.15329 (19)	0.43556 (8)	0.0191 (4)
H5	0.7319	1.1952	0.4380	0.023*
C6	0.6335 (2)	1.04440 (19)	0.40462 (7)	0.0171 (4)
H6	0.7093	1.0112	0.3866	0.021*
C7	0.5517 (2)	1.3217 (2)	0.49545 (8)	0.0247 (5)
H7A	0.4636	1.3631	0.4993	0.037*

H7B	0.6142	1.3815	0.4785	0.037*
H7C	0.5868	1.2985	0.5296	0.037*
C8	0.45172 (19)	0.45546 (18)	0.34029 (7)	0.0137 (4)
C9	0.45326 (19)	0.31071 (18)	0.33494 (7)	0.0137 (4)
H9A	0.5472	0.2798	0.3315	0.016*
H9B	0.4132	0.2705	0.3660	0.016*
C10	0.3811 (2)	0.13288 (17)	0.27813 (7)	0.0142 (4)
H10A	0.3570	0.0845	0.3098	0.017*
H10B	0.4738	0.1085	0.2680	0.017*
C11	0.28461 (19)	0.09925 (18)	0.23534 (7)	0.0132 (4)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0104 (2)	0.0118 (2)	0.0153 (2)	-0.00084 (17)	-0.00072 (16)	0.00057 (16)
O1	0.0240 (8)	0.0125 (7)	0.0221 (7)	-0.0013 (6)	-0.0072 (6)	0.0031 (5)
O2	0.0164 (7)	0.0158 (7)	0.0168 (7)	-0.0021 (6)	0.0028 (5)	-0.0016 (5)
O3	0.0109 (7)	0.0226 (8)	0.0233 (7)	-0.0016 (6)	-0.0006 (6)	-0.0024 (5)
O4	0.0169 (7)	0.0148 (7)	0.0197 (7)	0.0007 (6)	-0.0025 (5)	0.0007 (5)
O5	0.0293 (9)	0.0111 (7)	0.0192 (7)	-0.0007 (6)	-0.0092 (6)	-0.0014 (5)
O6	0.0156 (7)	0.0155 (7)	0.0217 (7)	-0.0016 (6)	-0.0030 (5)	0.0033 (5)
O7	0.0134 (7)	0.0143 (7)	0.0211 (7)	-0.0023 (6)	-0.0006 (6)	-0.0043 (5)
N1	0.0123 (8)	0.0109 (8)	0.0154 (8)	0.0011 (6)	-0.0002 (6)	0.0000 (6)
C1	0.0144 (9)	0.0131 (9)	0.0149 (8)	-0.0001 (7)	-0.0003 (7)	0.0010 (7)
C2	0.0148 (10)	0.0198 (10)	0.0215 (9)	-0.0019 (8)	0.0032 (8)	0.0004 (8)
C3	0.0190 (11)	0.0217 (10)	0.0228 (10)	0.0025 (8)	0.0066 (8)	-0.0020 (8)
C4	0.0237 (11)	0.0162 (10)	0.0150 (9)	0.0007 (8)	-0.0001 (8)	0.0004 (7)
C5	0.0145 (10)	0.0225 (10)	0.0202 (9)	-0.0045 (8)	0.0002 (8)	-0.0023 (8)
C6	0.0130 (9)	0.0199 (10)	0.0183 (9)	0.0004 (8)	0.0014 (7)	-0.0027 (7)
C7	0.0276 (12)	0.0219 (11)	0.0247 (10)	-0.0011 (9)	0.0010 (9)	-0.0067 (8)
C8	0.0102 (9)	0.0150 (9)	0.0161 (8)	-0.0005 (7)	0.0005 (7)	0.0004 (7)
C9	0.0125 (9)	0.0133 (9)	0.0152 (9)	0.0002 (7)	-0.0026 (7)	0.0000 (7)
C10	0.0132 (9)	0.0095 (8)	0.0200 (9)	0.0004 (7)	-0.0017 (7)	-0.0009 (7)
C11	0.0108 (9)	0.0132 (9)	0.0157 (8)	-0.0019 (7)	0.0028 (7)	0.0012 (7)

Geometric parameters (\AA , ^\circ)

S1—O3	1.4478 (14)	C2—H2	0.9500
S1—O1	1.4547 (14)	C3—C4	1.386 (3)
S1—O2	1.4675 (13)	C3—H3	0.9500
S1—C1	1.768 (2)	C4—C5	1.394 (3)
O4—C8	1.207 (2)	C4—C7	1.507 (3)
O5—C8	1.320 (2)	C5—C6	1.390 (3)
O5—H5o	0.84 (1)	C5—H5	0.9500
O6—C11	1.205 (2)	C6—H6	0.9500
O7—C11	1.328 (2)	C7—H7A	0.9800
O7—H7o	0.84 (1)	C7—H7B	0.9800
N1—C9	1.484 (2)	C7—H7C	0.9800

N1—C10	1.486 (2)	C8—C9	1.507 (3)
N1—H1n1	0.88 (1)	C9—H9A	0.9900
N1—H1n2	0.88 (1)	C9—H9B	0.9900
C1—C2	1.385 (3)	C10—C11	1.505 (3)
C1—C6	1.394 (3)	C10—H10A	0.9900
C2—C3	1.391 (3)	C10—H10B	0.9900
O3—S1—O1	114.00 (9)	C4—C5—H5	119.7
O3—S1—O2	112.28 (8)	C5—C6—C1	119.93 (18)
O1—S1—O2	111.72 (8)	C5—C6—H6	120.0
O3—S1—C1	106.53 (9)	C1—C6—H6	120.0
O1—S1—C1	105.95 (9)	C4—C7—H7A	109.5
O2—S1—C1	105.63 (8)	C4—C7—H7B	109.5
C8—O5—H5O	108.2 (18)	H7A—C7—H7B	109.5
C11—O7—H7O	110 (2)	C4—C7—H7C	109.5
C9—N1—C10	112.10 (14)	H7A—C7—H7C	109.5
C9—N1—H1N1	111.7 (17)	H7B—C7—H7C	109.5
C10—N1—H1N1	109.4 (16)	O4—C8—O5	125.13 (18)
C9—N1—H1N2	110.2 (14)	O4—C8—C9	123.20 (17)
C10—N1—H1N2	108.3 (15)	O5—C8—C9	111.66 (15)
H1N1—N1—H1N2	105 (2)	N1—C9—C8	109.01 (15)
C2—C1—C6	119.85 (18)	N1—C9—H9A	109.9
C2—C1—S1	120.41 (15)	C8—C9—H9A	109.9
C6—C1—S1	119.73 (15)	N1—C9—H9B	109.9
C1—C2—C3	119.53 (19)	C8—C9—H9B	109.9
C1—C2—H2	120.2	H9A—C9—H9B	108.3
C3—C2—H2	120.2	N1—C10—C11	109.43 (15)
C4—C3—C2	121.46 (18)	N1—C10—H10A	109.8
C4—C3—H3	119.3	C11—C10—H10A	109.8
C2—C3—H3	119.3	N1—C10—H10B	109.8
C3—C4—C5	118.49 (18)	C11—C10—H10B	109.8
C3—C4—C7	121.06 (19)	H10A—C10—H10B	108.2
C5—C4—C7	120.4 (2)	O6—C11—O7	125.83 (17)
C6—C5—C4	120.68 (19)	O6—C11—C10	123.36 (17)
C6—C5—H5	119.7	O7—C11—C10	110.79 (16)
O3—S1—C1—C2	16.35 (19)	C3—C4—C5—C6	0.9 (3)
O1—S1—C1—C2	-105.40 (17)	C7—C4—C5—C6	-178.42 (18)
O2—S1—C1—C2	135.94 (16)	C4—C5—C6—C1	1.2 (3)
O3—S1—C1—C6	-163.36 (15)	C2—C1—C6—C5	-2.3 (3)
O1—S1—C1—C6	74.89 (17)	S1—C1—C6—C5	177.39 (15)
O2—S1—C1—C6	-43.77 (18)	C10—N1—C9—C8	175.26 (15)
C6—C1—C2—C3	1.3 (3)	O4—C8—C9—N1	-5.4 (3)
S1—C1—C2—C3	-178.42 (15)	O5—C8—C9—N1	175.76 (15)
C1—C2—C3—C4	0.9 (3)	C9—N1—C10—C11	172.17 (15)
C2—C3—C4—C5	-1.9 (3)	N1—C10—C11—O6	-6.8 (3)
C2—C3—C4—C7	177.37 (19)	N1—C10—C11—O7	174.56 (14)

Hydrogen-bond geometry (Å, °)

<i>D—H···A</i>	<i>D—H</i>	<i>H···A</i>	<i>D···A</i>	<i>D—H···A</i>
N1—H1N1···O2 ⁱ	0.88 (1)	2.02 (1)	2.885 (2)	167 (2)
N1—H1N2···O3 ⁱⁱ	0.88 (1)	2.06 (2)	2.792 (2)	140 (2)
O5—H5O···O1	0.84 (1)	1.79 (1)	2.607 (2)	164 (3)
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