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Key indicators

Single-crystal X-ray study T = 295 KMean $\sigma(\text{C}-\text{C}) = 0.002 \text{ Å}$ R factor = 0.028 wR factor = 0.084 Data-to-parameter ratio = 11.9

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3-Ammonio-4-methoxybenzenesulfonate

3–Amino-4-methoxybenzenesulfonic acid exists in the solid state in the zwitterionic form as 3-ammonio-4-methoxybenzenesulfonate, $C_7H_9NO_4S$. The zwitterions are linked by the ammonium H atoms into a layer structure.

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Comment

3-Amino-4-hydroxybezenesulfonic acid, like a number of arenesulfonic acids, adopts a layer structure in which the aromatic rings stack into sheets; the sulfonate groups are located on the top and bottom of the layers (Gunderman & Squattrito, 1996). The title compound, (I), which has a methoxy substituent in place of the hydroxy substituent, also adopts a layer structure in which the zwitterions are linked to each other by hydrogen bonds (Fig. 1). The ammonium unit serves as the donor site for three hydrogen bonds; however, two of the three sulfonate O atoms are engaged in hydrogen bonding (Table 2). The negative charge appears to be delocalized over only two of the three O atoms, as the O atom that is not engaged in the interactions is nearest to the S atom.



Experimental

The title compound was recovered unchanged in an unsuccessful reaction with calcium nitrate hexahydrate. The calcium salt (0.82 g, 3 mmol) was mixed with methoxybenzenesulfonic acid (1.22 g, 6 mmol) in water. Colorless prismatic crystals separated from the solution after several days. Analysis calculated for $C_7H_9NO_4S$: C 41.37, H 4.46, N 6.89%; found C 41.33, H 4.48, N 6.87%.

Crystal data C7H9NO4S Z = 2 $M_r = 203.21$ $D_x = 1.678 \text{ Mg m}^{-3}$ Triclinic, $P\overline{1}$ Mo $K\alpha$ radiation Cell parameters from 3869 a = 7.321(2) Å b = 8.234(2) Å reflections $\theta = 3.1-27.5^{\circ}$ c = 8.354 (2) Å $\mu = 0.38 \text{ mm}^{-1}$ $\alpha = 62.05 (3)^{\circ}$ $\beta = 65.34 (3)^{\circ}$ T = 295 (2) K $\gamma = 74.04 \ (3)^{\circ}$ Prism, colorless $V = 402.3 (1) \text{ Å}^3$ $0.34 \times 0.26 \times 0.19 \text{ mm}$

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organic papers

Data collection

Rigaku R-AXIS RAPID diffractometer ω scans Absorption correction: multi-scan (*ABSCOR*; Higashi, 1995) *T*_{min} = 0.810, *T*_{max} = 0.931 3979 measured reflections

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.028$ $wR(F^2) = 0.084$ S = 1.041830 reflections 154 parameters All H-atom parameters refined 1830 independent reflections 1713 reflections with $I > 2\sigma(I)$ $R_{int} = 0.011$ $\theta_{max} = 27.5^{\circ}$ $h = -9 \rightarrow 8$ $k = -10 \rightarrow 10$ $l = -10 \rightarrow 10$

 $w = 1/[\sigma^{2}(F_{o}^{2}) + (0.053P)^{2} + 0.1821P]$ where $P = (F_{o}^{2} + 2F_{c}^{2})/3$ $(\Delta/\sigma)_{\text{max}} = 0.001$ $\Delta\rho_{\text{max}} = 0.33 \text{ e} \text{ Å}^{-3}$ $\Delta\rho_{\text{min}} = -0.38 \text{ e} \text{ Å}^{-3}$

Table 1

Selected geometric parameters (Å, °).

S1-O1	1.465 (1)	O4-C4	1.354 (2)
S1-O2	1.464 (1)	O4-C7	1.434 (2)
S1-O3	1.445 (1)	N1-C3	1.457 (2)
S1-C1	1.771 (2)		
O1-S1-O2	111.1 (1)	C2-C1-C6	120.4 (1)
O1-S1-O3	114.4 (1)	C2-C1-S1	120.0 (1)
O2-S1-O3	112.6(1)	C6-C1-S1	119.5 (1)
C1-S1-O1	105.0(1)	C2-C3-N1	120.4 (1)
C1-S1-O2	105.4 (1)	C4-C3-N1	117.9 (1)
C1-S1-O3	107.6(1)	O4-C4-C5	125.9 (1)
C4-O4-C7	117.8 (1)	O4-C4-C3	115.1 (1)

Table 2Hydrogen-bonding geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1-H1n1\cdotsO1^{i}$ $N1-H1n2\cdotsO2^{ii}$	0.86(1) 0.86(1)	1.98 (1) 1.93 (1)	2.829 (2) 2.770 (2)	171 (2) 169 (2)
$N1-H1n3\cdotsO1^{iii}$	0.86 (1)	2.10(1)	2.878 (2)	151 (2)

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

All H atoms were refined with distance restraints of N-H = 0.85 (1) Å and C-H = 0.95 (1) Å.



Figure 1

ORTEPII (Johnson, 1976) plot of (I). Displacement ellipsoids are drawn at the 70% probability level and H atoms are drawn as spheres of arbitrary radii.

Data collection: *RAPID-AUTO* (Rigaku, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (Rigaku/MSC, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); 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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