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Ethane-1,2-diaminium 4,4'-sulfonyldibenzoate

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Key indicators: single-crystal X-ray study; T = 200 K; mean σ (C–C) = 0.003 Å; R factor = 0.034; wR factor = 0.098; data-to-parameter ratio = 14.1.

In the title salt, $C_2H_{10}N_2^{2+}C_{14}H_8O_6S^{2-}$, both the ethylenediaminium cations and the 4,4'-sulfonyldibenzoate dianions have crystallographic twofold rotational symmetry. They are interlinked by aminium N-H···O_{carboxylate} hydrogen-bonding associations, giving sheets parallel to (101) and are further linked along [010], forming a three-dimensional structure.

Related literature

For the structure of 4,4'-sulfonyldibenzoic acid, see: Lian et al. (2007). For the structures of some metal complexes of the acid, see: Bannerjee et al. (2009); Jiao (2010); Pan et al. (2007); Wu et al. (2007); Zhuang & Jin (2007).



Experimental

Crystal data

 $C_2H_{10}N_2^{2+}C_{14}H_8O_6S^{2-}$ $M_{\rm w} = 366.39$ Monoclinic, P2/c a = 15.2860 (8) Å b = 4.8436 (2) Å c = 11.9803 (6) Å $\beta = 111.812 \ (6)^{\circ}$

Data collection

Oxford Diffraction Gemini-S CCD detector diffractometer Absorption correction: multi-scan (CrysAlis PRO; Oxford Diffraction, 2010) $T_{\min} = 0.98, T_{\max} = 0.99$

V = 823.51 (8) Å³ Z = 2Mo $K\alpha$ radiation $\mu = 0.23 \text{ mm}^{-1}$ T = 200 K $0.35 \times 0.25 \times 0.08 \text{ mm}$

5062 measured reflections 1607 independent reflections 1290 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.024$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.034$	114 parameters
$wR(F^2) = 0.098$	H-atom parameters constrained
S = 1.05	$\Delta \rho_{\rm max} = 0.35 \ {\rm e} \ {\rm \AA}^{-3}$
1607 reflections	$\Delta \rho_{\rm min} = -0.25 \ {\rm e} \ {\rm \AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

$D - H \cdots A$	D-H	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1A - H11A \cdots O42^{i}$	0.89	1.87	2.760 (2)	174
$N1A - H12A \cdots O41$ $N1A - H13A \cdots O42$	0.89	1.88	2.740 (2) 2.798 (2)	163

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

Data collection: CrysAlis PRO (Oxford Diffraction, 2010); cell refinement: CrysAlis PRO; data reduction: CrysAlis PRO; program(s) used to solve structure: SIR92 (Altomare et al., 1994); program(s) used to refine structure: SHELXL97 (Sheldrick, 2008) within WinGX (Farrugia, 1999); molecular graphics: PLATON (Spek, 2009); software used to prepare material for publication: PLATON.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: LH5347).

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supplementary materials

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Ethane-1,2-diaminium 4,4'-sulfonyldibenzoate

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Comment

The structure of the diprotic acid 4,4'-sulfonyldibenzoic acid (SDBA) is known (Lian *et al.*, 2007) and although some metal complexes with SDBA alone have been reported, *e.g.* with Li (Bannerjee *et al.*, 2009), Zn (Pan *et al.*, 2007; Zhuang & Jin, 2007) and Cd (Jiao, 2010), most structures have been with mixed ligands including this acid (Wu *et al.*, 2007). No structures of compounds of SDBA with Lewis bases are known. Our 1:1 stoichiometric reaction of this acid with ethylenediamine gave the the title compound $C_2H_{10}N_2^{2+}C_{14}H_8O_6S^{2-}$, and the structure is reported here. In this structure (Fig. 1), both the ethylenediaminium cations and the 4,4'-sulfonyldibenzoate dianions have crystallographic twofold rotational symmetry. In contrast, the two substituted ring systems of the parent molecule are mirror related (Lian *et al.*, 2007). With the present salt, the central C1—S1—C1ⁱ bond angle is 104.90 (8)Å [for symmetry code (i), see Fig. 1], while the carboxyl group (defined by atoms C4–C41–O41–O42) lies slightly out of the plane of the benzene ring [dihedral angle 19.31 (9)°]. The ethylenediamine cation is essentially planar [torsion angle N1A—C1A—C1Aⁱⁱ—N1Aⁱⁱ, 171.97 (14)°]. For symmetry code (ii), see Also Fig. 1.

Intermolecular cation aminium $N - H \cdots O_{carboxyl}$ hydrogen bonds (Table 1) interlink the SDBA dianions into sheets lying in the (101) planes, as well as down the *b* axis, forming a three-dimensional structure (Fig. 2). The sulfonyl O atoms are involved in inter-species C - H \cdots O associations [C2 - H2 \cdots O1^v, 3.200 (2) Å: symmetry code (v) -*x* + 2, -*y* + 1, -*z* + 1]. as well as in S - O \cdots Cg interactions [minimum S1 - O1 \cdots Cg(ring C1^{viii} - C6^{viii}) = 3.5409 (13) Å; S - O \cdots Cg angle, 90.61 (5)°: symmetry code (viii) -*x* + 2, *y* + 1, -*z* + 3/2].

Experimental

The title compound was synthesized by heating together for 10 min under reflux, 1 mmol quantities of 4,4'-sulfonyldibenzoic acid and ethylenediamine in 50 ml of 50% ethanol–water. After evaporation of the solvent the non-crystalline product was recrystallized from a 50% methanol–isopropyl alcohol solution giving thin colourless crystal plates from which a specimen was cleaved for the X-ray analysis.

Refinement

The aminium H atoms were located by difference Fourier methods and their positional and isotropic displacement parameters were initially refined but finally were allowed to ride on the N atom with $U_{iso}(H) = 1.2U_{eq}(N)$. Other H atoms were included at calculated positions with C—H (aromatic) = 0.93 Å or C—H (methyl) = 0.97 Å] and also treated as riding, with $U_{iso}(H) = 1.2U_{eq}C(aromatic)$ or $1.5U_{eq}C(methylene)$.

Figures



Fig. 1. Molecular conformation and atom-numbering scheme for the title compound, with the inter-species hydrogen bond shown as a dashed line and with non-H atoms shown as 50% probability displacement ellipsoids. Both the dication and the dianion have twofold rotational symmetry [symmetry codes (i) -x + 2, y, -z + 3/2; (ii) -x + 1, y, -z + 1/2].

Fig. 2. A perspective view of the three-dimensional structure looking along the b axial direction, showing hydrogen-bonding associations as dashed lines. Carbon-bound H atoms are omitted. For symmetry codes, see Table 1 and Fig. 1.

Ethane-1,2-diaminium 4,4'-sulfonyldibenzoate

Crystal data

 $C_2H_{10}N_2^{2+}C_{14}H_8O_6S^{2-}$ F(000) = 384 $M_r = 366.39$ $D_{\rm x} = 1.478 \ {\rm Mg \ m}^{-3}$ Mo *K* α radiation, $\lambda = 0.71073$ Å Monoclinic, P2/c Hall symbol: -P 2yc Cell parameters from 2936 reflections $\theta = 3.5 - 28.6^{\circ}$ a = 15.2860 (8) Å b = 4.8436 (2) Å $\mu = 0.23 \text{ mm}^{-1}$ T = 200 Kc = 11.9803 (6) Å $\beta = 111.812 \ (6)^{\circ}$ Plate, colourless $0.35 \times 0.25 \times 0.08 \text{ mm}$ V = 823.51 (8) Å³ Z = 2

Data collection

Oxford Diffraction Gemini-S CCD detector diffractometer Radiation source: Enhance (Mo) X-ray source graphite Detector resolution: 16.077 pixels mm⁻¹ ω scans Absorption correction: multi-scan (CrysAlis PRO; Oxford Diffraction, 2010) $T_{\min} = 0.98, T_{\max} = 0.99$ 5062 measured reflections

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.034$ $wR(F^2) = 0.098$

1607 independent reflections 1290 reflections with $I > 2\sigma(I)$ $R_{\rm int} = 0.024$ $\theta_{\text{max}} = 26.0^{\circ}, \ \theta_{\text{min}} = 3.4^{\circ}$ $h = -18 \rightarrow 18$ $k = -5 \rightarrow 5$ $l = -14 \rightarrow 14$

Primary atom site location: structure-invariant direct methods Secondary atom site location: difference Fourier map Hydrogen site location: inferred from neighbouring sites H-atom parameters constrained

<i>S</i> = 1.05	$w = 1/[\sigma^2(F_0^2) + (0.0627P)^2]$ where $P = (F_0^2 + 2F_c^2)/3$
1607 reflections	$(\Delta/\sigma)_{\rm max} < 0.001$
114 parameters	$\Delta \rho_{max} = 0.35 \text{ e } \text{\AA}^{-3}$
0 restraints	$\Delta \rho_{\rm min} = -0.25 \ e \ {\rm \AA}^{-3}$

Special details

Geometry. Bond distances, angles *etc.* have been calculated using the rounded fractional coordinates. All su's are estimated from the variances of the (full) variance-covariance matrix. The cell e.s.d.'s are taken into account in the estimation of distances, angles and torsion angles

Refinement. Refinement of F^2 against ALL reflections. The weighted *R*-factor *wR* and goodness of fit *S* are based on F^2 , conventional *R*-factors *R* are based on *F*, with *F* set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating *R*-factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. *R*-factors based on F^2 are statistically about twice as large as those based on *F*, and *R*- factors based on ALL data will be even larger.

	x	У	Ζ	$U_{\rm iso}*/U_{\rm eq}$
S1	1.00000	0.61975 (12)	0.75000	0.0180 (2)
O1	1.03086 (9)	0.7681 (3)	0.66702 (11)	0.0249 (4)
O41	0.63470 (9)	-0.2635 (3)	0.51471 (11)	0.0301 (4)
O42	0.66855 (9)	-0.1734 (3)	0.35316 (10)	0.0243 (4)
C1	0.90679 (11)	0.3967 (3)	0.66620 (15)	0.0175 (5)
C2	0.90534 (12)	0.2888 (4)	0.55790 (16)	0.0225 (5)
C3	0.83192 (12)	0.1147 (4)	0.49238 (15)	0.0221 (5)
C4	0.76174 (11)	0.0437 (4)	0.53517 (14)	0.0176 (5)
C5	0.76491 (13)	0.1538 (4)	0.64426 (16)	0.0216 (5)
C6	0.83708 (12)	0.3320 (4)	0.71004 (16)	0.0221 (5)
C41	0.68270 (12)	-0.1474 (3)	0.46402 (15)	0.0192 (5)
N1A	0.61734 (10)	0.3312 (3)	0.23264 (13)	0.0242 (5)
C1A	0.51341 (13)	0.3110 (4)	0.19485 (16)	0.0273 (6)
H2	0.95280	0.33250	0.52980	0.0270*
Н3	0.82960	0.04480	0.41900	0.0260*
H5	0.71830	0.10750	0.67330	0.0260*
H6	0.83850	0.40660	0.78220	0.0260*
H11A	0.63740	0.48570	0.27460	0.0290*
H12A	0.63270	0.33380	0.16790	0.0290*
H13A	0.64420	0.18630	0.27810	0.0290*
H14A	0.48380	0.46620	0.14350	0.0330*
H15A	0.49130	0.14250	0.14940	0.0330*

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

Atomic displacement parameters (A ⁺	A* ,	1-))
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	U^{11}	U ²²	U^{33}	U^{12}	U^{13}	U^{23}
S1	0.0190 (3)	0.0165 (3)	0.0174 (3)	0.0000	0.0055 (2)	0.0000

supplementary materials

O1	0.0270 (7)	0.0217 (7)	0.0255 (7)	-0.0024 (5)	0.0091 (6)	0.0053 (6)
O41	0.0338 (8)	0.0381 (8)	0.0204 (7)	-0.0146 (6)	0.0123 (6)	-0.0022 (6)
O42	0.0310 (7)	0.0265 (7)	0.0153 (6)	-0.0061 (5)	0.0086 (5)	-0.0033 (5)
C1	0.0170 (8)	0.0169 (9)	0.0167 (8)	0.0018 (6)	0.0041 (7)	0.0021 (7)
C2	0.0214 (9)	0.0292 (10)	0.0195 (9)	-0.0025 (7)	0.0107 (7)	-0.0014 (8)
C3	0.0257 (9)	0.0265 (10)	0.0158 (9)	-0.0013 (8)	0.0098 (7)	-0.0031 (7)
C4	0.0177 (8)	0.0187 (9)	0.0148 (8)	0.0021 (7)	0.0041 (7)	0.0028 (7)
C5	0.0237 (9)	0.0244 (10)	0.0214 (9)	-0.0038 (7)	0.0140 (8)	-0.0008 (7)
C6	0.0268 (9)	0.0244 (10)	0.0169 (9)	-0.0014 (8)	0.0103 (8)	-0.0037 (7)
C41	0.0215 (9)	0.0185 (9)	0.0179 (9)	0.0027 (7)	0.0077 (7)	0.0027 (7)
N1A	0.0292 (9)	0.0254 (8)	0.0189 (8)	-0.0004 (7)	0.0099 (7)	-0.0003 (7)
C1A	0.0245 (10)	0.0361 (11)	0.0209 (10)	0.0020 (8)	0.0080 (8)	-0.0024 (8)

Geometric parameters (Å, °)

S1—O1	1.4407 (14)	C2—C3	1.390 (3)
S1—C1	1.7726 (17)	C3—C4	1.393 (3)
\$1-01 ⁱ	1.4407 (14)	C4—C41	1.509 (2)
81—C1 ⁱ	1.7726 (17)	C4—C5	1.396 (2)
O41—C41	1.247 (2)	C5—C6	1.392 (3)
O42—C41	1.270 (2)	C2—H2	0.9300
N1A—C1A	1.485 (3)	С3—Н3	0.9300
N1A—H13A	0.8900	С5—Н5	0.9300
N1A—H11A	0.8900	С6—Н6	0.9300
N1A—H12A	0.8900	C1A—C1A ⁱⁱ	1.522 (3)
C1—C2	1.391 (2)	C1A—H14A	0.9700
C1—C6	1.388 (3)	C1A—H15A	0.9700
01—S1—C1	108.27 (8)	C4—C5—C6	120.77 (18)
01—S1—O1 ⁱ	120.17 (9)	C1—C6—C5	118.97 (17)
01—S1—C1 ⁱ	107.12 (8)	O42—C41—C4	116.17 (16)
01 ⁱ —S1—C1	107.12 (8)	O41—C41—O42	124.21 (16)
C1—S1—C1 ⁱ	104.90 (8)	O41—C41—C4	119.61 (15)
O1 ⁱ —S1—C1 ⁱ	108.27 (8)	C3—C2—H2	121.00
C1A—N1A—H11A	109.00	C1—C2—H2	120.00
C1A—N1A—H12A	110.00	С2—С3—Н3	120.00
C1A—N1A—H13A	109.00	С4—С3—Н3	120.00
H11A—N1A—H12A	109.00	C4—C5—H5	120.00
H11A—N1A—H13A	109.00	С6—С5—Н5	120.00
H12A—N1A—H13A	109.00	С5—С6—Н6	120.00
C2—C1—C6	121.32 (16)	С1—С6—Н6	121.00
S1—C1—C2	119.21 (14)	N1A—C1A—C1A ⁱⁱ	109.76 (15)
S1—C1—C6	119.47 (13)	N1A—C1A—H14A	110.00
C1—C2—C3	118.96 (17)	N1A—C1A—H15A	110.00
C2—C3—C4	120.83 (16)	H14A—C1A—H15A	108.00
C3—C4—C41	120.53 (15)	C1A ⁱⁱ —C1A—H14A	110.00
C3—C4—C5	119.13 (17)	C1A ⁱⁱ —C1A—H15A	110.00
C5—C4—C41	120.35 (16)		

01 - S1 - C1 - C2 01 - S1 - C1 - C6	30.61 (16) -149 72 (14)	C2-C3-C4-C5 C2-C3-C4-C41	1.2 (3) -179 47 (17)
01^{i} S1 - C1 - C2	161.56 (14)	C3-C4-C5-C6	-0.1 (3)
O1 ⁱ —S1—C1—C6	-18.77 (16)	C41—C4—C5—C6	-179.47 (17)
C1 ⁱ —S1—C1—C2	-83.51 (15)	C3—C4—C41—O41	162.10 (17)
C1 ⁱ —S1—C1—C6	96.15 (15)	C3—C4—C41—O42	-19.4 (2)
S1—C1—C2—C3	-179.80 (14)	C5—C4—C41—O41	-18.5 (3)
C6—C1—C2—C3	0.5 (3)	C5—C4—C41—O42	159.93 (17)
S1—C1—C6—C5	-179.17 (14)	C4—C5—C6—C1	-0.7 (3)
C2-C1-C6-C5	0.5 (3)	N1A—C1A—C1A ⁱⁱ —N1A ⁱⁱ	171.97 (14)
C1—C2—C3—C4	-1.4 (3)		
Symmetry codes: (i) $-x+2$, y, $-z+3/2$; (i	i) $-x+1$, y , $-z+1/2$.		

Hydrogen-bond geometry (Å, °)

D—H···A	<i>D</i> —Н	$H \cdots A$	$D \cdots A$	D—H···A
N1A—H11A···O42 ⁱⁱⁱ	0.89	1.87	2.760 (2)	174
N1A—H12A···O41 ^{iv}	0.89	1.88	2.740 (2)	163
N1A—H13A…O42	0.89	1.93	2.798 (2)	164
C2— $H2$ ···O1 ^v	0.93	2.51	3.200 (2)	131
C5—H5···O42 ^{vi}	0.93	2.56	3.344 (2)	143
C6—H6···O1 ⁱ	0.93	2.55	2.915 (2)	104
C1A—H14A···O41 ^{vii}	0.97	2.46	3.384 (2)	160
Symmetry codes: (iii) $x, y+1, z$; (iv) $x, -y, z-1/2$; (v)	-x+2, -y+1, -z+1; (x)	<i>x</i> , − <i>y</i> , <i>z</i> +1/2; (i) − <i>z</i>	x+2, y, -z+3/2; (vii)	-x+1, y+1, -z+1/2.

Fig. 1





Fig. 2