C(16)—C(17)	1.505 (7)	C(16)—C(18)	1.505 (4)	
C(19)—C(20)	1.521 (5)	C(20)—C(21)	1.520 (5)	
C(21)—C(22)	1.511 (6)			
C(1)—O(2)—C(5)	109.1 (2)	C(14)—O(3)—C(18)	110.1 (2)	
C(10)NC(13)	109.3 (2)	C(10)NC(19)	123.1 (2)	
C(13—N—C(19)	115.9 (2)	O(1)-C(1)-O(2)	121.6 (3)	
O(1) - C(1) - C(2)	128.3 (3)	O(2) - C(1) - C(2)	110.1 (3)	
C(1) - C(2) - C(3)	112.5 (3)	C(1)—C(2)—C(4)	101.4 (2)	
C(3)C(2)C(4)	120.6 (2)	C(2)—C(4)—C(5)	100.7 (2)	
C(2)—C(4)—C(11)	110.9 (2)	C(5)—C(4)—C(11)	113.7 (2)	
O(2)C(5)C(4)	103.3 (2)	O(2)—C(5)—C(6)	109.9 (2)	
C(4)—C(5)—C(6)	118.1 (2)	C(5)—C(6)—C(7)	111.4 (3)	
C(5)—C(6)—C(9)	111.1 (2)	C(7)—C(6)—C(9)	112.3 (3)	
C(6)—C(7)—C(8)	114.8 (3)	C(6)—C(9)—C(10)	113.5 (2)	
C(6)—C(9)—C(22)	116.4 (3)	C(10)—C(9)—C(22)	112.6 (2)	
N-C(10)-C(9)	120.2 (2)	N-C(10)-C(11)	105.4 (2)	
C(9)—C(10)—C(11)	113.5 (2)	C(4)—C(11)—C(10)	115.5 (2)	
C(4)—C(11)—C(12)	111.8 (2)	C(10)—C(11)—C(12)	101.8 (2)	
C(11)—C(12)—C(13)	103.2 (2)	N-C(13)-C(12)	104.4 (2)	
N-C(13)-C(14)	114.3 (2)	C(12)—C(13)—C(14)	109.6 (2)	
O(3)-C(14)-C(13)	110.1 (2)	O(3)-C(14)-C(15)	103.5 (2)	
C(13)—C(14)—C(15)	115.3 (2)	C(14)-C(15)-C(16)	104.0 (3)	
C(15)-C(16)-C(17)	116.5 (4)	C(15)-C(16)-C(18)	102.5 (2)	
C(17)—C(16)—C(18)	114.2 (3)	O(3)-C(18)-O(4)	120.7 (3)	
O(3)-C(18)-C(16)	110.5 (3)	O(4)-C(18)-C(16)	128.8 (3)	
N-C(19)-C(20)	114.1 (3)	C(19)—C(20)—C(21)	114.5 (3)	
C(20)—C(21)—C(22)	115.4 (3)	C(9)—C(22)—C(21)	116.8 (3)	
C(10)NC(19)—C(20)	64.4 (5)	
C(19)—C(20)—	-C(21)-C(22)	63.8 (6	5)	
C(21)—C(22)—	-C(9)-C(10)	77.7 (5)	
C(9)—C(10)—	N—C(19)	-1.3 (5	5)	
NC(19)C(2	20)—C(21)	-80.1 (5)		
C(20)—C(21)—	-C(22)-C(9)	-64.7 (5)		
C(22)—C(9)—	C(10)—N	-60.1 (5)	

The structure was solved by direct methods using *SHELXS86* (Sheldrick, 1985). H atoms were located in a difference Fourier synthesis using *Xtal* (Stewart & Hall, 1983).

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Lists of structure factors, anisotropic displacement parameters and H-atom coordinates have been deposited with the IUCr (Reference: KA1019). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Interactions Between Sulfonated Azo Dyes and Biomolecules: Orange G/Adenine and Orange G/Cytosine Salts

WILLIAM H. OJALA AND WILLIAM B. GLEASON

Biomedical Engineering Center, Department of Laboratory Medicine and Pathology, University of Minnesota, Minneapolis, MN 55455, USA

TIMOTHY I. RICHARDSON AND REX E. LOVRIEN

Department of Biochemistry, University of Minnesota, St. Paul, MN 55108, USA

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Abstract

The disulfonated azo dye Orange G [the disodium] salt of 7-hydroxy-8-(phenylazo)-1,3-naphthalenedisulfonic acid] forms salts with adenine and cytosine on co-crystallization from aqueous HCl. The 1:2 dye:adenine crystal, $2C_5H_6N_5^+.C_{16}H_{10}N_2O_7S_2^{2-}$.-5H₂O, is a pentahydrate and the 1:2 dye:cytosine crystal, $2C_4H_6N_3O^+.C_{16}H_{10}N_2O_7S_2^{2-}.H_2O$, is a monohydrate. In the solid state the dye is found to exist predominantly as the hydrazo, rather than the azo, tautomer. In both structures, one of the protonated nucleotide bases approaches close to a sulfonate group of the dye in an 'edge-on' fashion. Molecules in the Orange G/adenine structure lie in segregated layers, but molecules in the Orange G/ cytosine structure lie in mixed stacks.

Comment

Interactions between biomolecules containing sulfate groups and biomolecules (such as proteins) capable of recognizing these groups are currently the object of intensive biomedical study. Although the focus of

such investigations is often on large sulfated molecules such as heparin, we are currently studying interactions between smaller molecules, particularly between sulfonated azo dyes and biomolecules, such as amino acids and nucleic acid bases, to determine whether the sulfonate group can serve as a mimic of the sulfate group. Sulfonated azo dyes have proved effective as co-precipitating and co-crystallizing agents for peptides and other biomolecules (Conrov & Lovrien, 1992). In addition, these dyes are of special interest in their own right because their potential usefulness as pharmaceuticals, especially as antiviral agents, has recently been recognized (Weaver, Pine, Anand, Bell & Aszalos, 1992). Understanding the biological activity of sulfated/sulfonated compounds and modeling their interactions with proteins or nucleic acids is greatly facilitated by the availability of structural information from X-ray crystallographic studies. Here we describe the structures of two crystalline dye/biomolecule salts grown from 0.2 N HCl solution, one consisting of the dye Orange G and adenine, the other consisting of Orange G and cytosine.





In each crystal structure the asymmetric unit was found to contain one dye molecule (as the dianion) and two protonated nucleotide base molecules (Figs. 1 and 2). Location of the corresponding H atoms in difference maps identified the protonation sites as the ring N atoms adjacent to the amino groups for both adenine and cytosine. The asymmetric unit of Orange G/adenine also contains five water molecules, while that of Orange G/cytosine contains only a single water molecule.

The geometry of the dye molecule in both structures is of interest in several respects. An azo compound with a suitably placed naphtholic hydroxyl group may exist as a hydrazone, as an azo compound, or as an equilibrium mixture of the two tautomers either in solution or in the solid state



Fig. 1. View of the Orange G/adenine asymmetric unit, showing atom numbering. For non-H atoms, 50% probability ellipsoids are shown.



Fig. 2. View of the Orange G/cytosine asymmetric unit, showing atom numbering. For non-H atoms, 50% probability ellipsoids are shown.

(Olivieri, Wilson, Paul & Curtin, 1989). In the crystal structures described here, the location of the H atom on N(2) rather than on O(7) and the lengths of the N(1)-N(2), N(1)-C(1), and C(2)-O(7) bonds indicate that here Orange G exists primarily as the hydrazo tautomer rather than as the azo tautomer. In both structures the S(1)-C(9) bond is significantly longer than the S(2)—C(7) bond, perhaps as a result of crowding by the bulky phenylazo substituent located near the S(1) sulfonate group. We have also observed this bond-length difference in the crystal structures of the diammonium, dilithium, magnesium, and calcium salts of Orange G (Ojala et al., 1994). In Orange G/adenine and Orange G/cytosine the conformations of the dye molecules resemble each other closely, with the phenylazo groups assuming almost identical orientations with respect to the naphthalene rings. Corresponding torsion angles describing the geometry of the phenylazo groups are very similar (Tables 3 and 4).

Noteworthy among the many hydrogen bonds in the structures (Table 5) are contacts between two of the O atoms from a given sulfonate group and two N-H groups from a neighboring protonated nucleotide base molecule. This two-pronged sulfonate-bridging contact, found in both crystal structures $[O(1) \cdots N(4) = 2.742 (4), O(2) \cdots N(3) = 2.809 (4) Å$ in Orange G/adenine; $O(2) \cdots N(6) =$ 2.857 (2), O(3)...N(7) = 2.740 (2) Å in Orange G/ cytosine], is suggestive of the type of contact expected between a sulfonate group (or sulfate group) and the guanidinium terminus of an arginine side chain, an interaction likely to occur between sulfonated (or sulfated) molecules and peptides. In both structures the interacting O-S-O and N-C-N groups approach each other in approximately the same plane [the angle between the planes O(1)-S(1)-O(2) and N(3)-C(17)-N(4) in Orange G/adenine is $18.3 (3)^{\circ}$; the angle between the planes O(2)-S(1)-O(3) and N(6)-C(21)-N(7) in Orange G/cytosine is $26.3 (2)^{\circ}$].

Inspection of the packing arrangements assumed in these two structures reveals a striking difference between adenine and cytosine molecules with respect to how they interact with and are packed with the molecules of Orange G. Orange G and adenine molecules do not mingle freely in the crystal but instead lie in segregated layers (Fig. 3). The water molecules present in the structure are found in the regions where these layers meet and into which the sulfonate groups project. In contrast, a neatly segregated layer structure is not found in Orange G/ cytosine. Instead the molecules lie in stacks in which cytosine and Orange G molecules overlap (Fig. 4). In these crystal structures, at least, the sulfonated azo dye appears to intercalate more readily with cytosine molecules than with adenine molecules.



Fig. 3. View of the Orange G/adenine molecular packing, showing alternating layers of dye and nucleotide base molecules. H atoms have been omitted for clarity.



Fig. 4. View of the Orange G/cytosine molecular packing, showing mixed stacks of dye and nucleotide base molecules. H atoms (except those on the cytosine amino groups) have been omitted for clarity.

Experimental

Co-precipitation and co-crystallization techniques used for growing these and similar co-crystals have been described previously (Conroy & Lovrien, 1992).

Orange G/Adenine

Crystal data	
$2C_5H_6N_5^+.C_{16}H_{10}N_2O_7S_2^{2-}$	Cu $K\alpha$ radiation
5H ₂ O	$\lambda = 1.54178 \text{ A}$
$M_r = 768.73$	Cell parameters from 25
Triclinic	reflections
PĪ	$\theta = 20.0 - 25.0^{\circ}$
a = 10.157 (4) Å	$\mu = 2.18 \text{ mm}^{-1}$
b = 10.277 (6) Å	T = 173 K
c = 17.551 (6) Å	Thin plate
$\alpha = 92.81 (4)^{\circ}$	$0.36 \times 0.12 \times 0.02 \text{ mm}$
$\beta = 93.07 (3)^{\circ}$	Orange
$\gamma = 118.00 (5)^{\circ}$	
$V = 1609 (3) Å^3$	
Z = 2	
$D_x = 1.586 \text{ Mg m}^{-3}$	

Data collection

 $R_{int} = 0.064$ Enraf-Nonius CAD-4 diffractometer $\theta_{\rm max} = 70.0^{\circ}$ $h = -12 \rightarrow 12$ $\omega/2\theta$ scans $k=-12 \rightarrow 12$ Absorption correction: $l = -17 \rightarrow 21$ empirical 3 standard reflections $T_{\min} = 0.80, T_{\max} = 1.00$ 9379 measured reflections frequency: 60 min intensity variation: none 6026 independent reflections 4785 observed reflections $[I > 3\sigma(I)]$

Refinement

Refinement on *F* R = 0.055wR = 0.075S = 2.68 $\begin{array}{l} \Delta \rho_{\rm max} = 0.65 \mbox{ e } {\rm \AA}^{-3} \\ \Delta \rho_{\rm min} = -0.65 \mbox{ e } {\rm \AA}^{-3} \\ {\rm Extinction \ correction:} \\ {\rm not \ applied} \end{array}$

4785 reflections	Atomic scattering factors	O(8W)	0.1040 (3)	0.5072 (3)	0.7217 (1)	3.5 (1)
496 narameters	from International Tables	O(9W)	-0.7767 (3)	-0.1925 (3)	0.6322 (1)	3.3 (1)
$w = 1/\sigma^2(F)$	for Y-ray Crystallography	O(10W)	-0.4160 (3)	0.5226 (3)	0.2358 (1)	3.19 (9)
$w = 1/0 (1^{\circ})$	$(1074 \text{ M}^{-1} \text{ M})$	O(11W)	-0.4090 (3)	0.2053 (4)	0.2273 (1)	3.9 (1)
$(\Delta/\sigma)_{\rm max} = 0.02$	(19/4, VOI. 1V)	O(12W)	-0.3694 (3)	0.9698 (4)	0.2140 (2)	5.1 (1)
		N(1)	0.2364 (3)	0.5229 (3)	0.0397 (1)	2.06 (8)
Orange G/Cytosine		N(2)	0.2844 (3)	0.4558 (3)	-0.0069(1)	2.27 (9)
Crustal data		N(3)	0.1363 (3)	0.3002 (4)	0.3296 (2)	2.5(1)
Crysiai aala		IN(4)	0.1948 (3)	0.5255 (5)	0.3979(2)	2.17(9)
$2C_4H_6N_3O^+$, $C_{16}H_{10}N_2O_7S_2^{2-}$.	Cu $K\alpha$ radiation	N(5)	0.1780 (3)	0.3369(3)	0.5500(1)	2.29 (9)
Но	$\lambda = 1.54178$ Å	N(7)	0.0800 (3)	0.3243(3) 0.1722(3)	0.3835(1) 0.4845(1)	2.30 (9)
M = 649.62	$C_{\rm cll} = 1.5 + 170 M$	N(8)	-0.4896(3)	0.1722(3) 0.4074(3)	0 3813 (2)	2.20 (9)
$M_r = 046.02$	Cell parameters nom 25	N(9)	-0.5629(3)	0.1589(3)	0.3524(1)	2.20 (9)
Triclinic	reflections	N(10)	-0.6615(3)	-0.0313(3)	0.4365 (2)	2.5 (1)
P1	$\theta = 20.0 - 25.0^{\circ}$	N(11)	-0.6711 (3)	0.0697 (3)	0.5632 (2)	2.5 (1)
a = 8.908 (2) Å	$\mu = 2.44 \text{ mm}^{-1}$	N(12)	-0.5781 (3)	0.3102 (3)	0.5430 (2)	2.5 (1)
b = 11.632 (4) Å	T = 173 K	C(1)	0.2009 (3)	0.6229 (4)	0.0135 (2)	2.1 (1)
c = 11.052 (4) R	T = 175 R Thin plate	C(2)	0.1967 (4)	0.6477 (4)	-0.0673 (2)	2.4 (1)
c = 13.097(2) A		C(3)	0.1313 (4)	0.7366 (4)	-0.0919 (2)	2.6 (1)
$\alpha = 90.17(3)^{\circ}$	$0.36 \times 0.16 \times 0.04 \text{ mm}$	C(4)	0.0760 (3)	0.7967 (4)	-0.0414 (2)	2.4 (1)
$\beta = 90.74 \ (2)^{\circ}$	Orange	C(5)	0.0830 (3)	0.7804 (4)	0.0401 (2)	2.0(1)
$\gamma = 110.96 (2)^{\circ}$		C(6)	0.0240 (3)	0.8482 (4)	0.0878 (2)	2.0(1)
$V = 1325 (1) Å^3$		C(7)	0.0366 (3)	0.8420 (4)	0.1662 (2)	1.84 (9)
7 - 2		C(8)	0.1156 (3)	0.7740 (4)	0.1966 (2)	1.9 (1)
L = 2		C(9)	0.1745 (3)	0.7046 (3)	0.1508 (2)	1.77 (9)
$D_x = 1.625 \text{ Mg m}^{-3}$		C(10)	0.1538 (3)	0.7004 (4)	0.0688 (2)	1.9 (1)
		$C(\Pi)$	0.3166 (3)	0.3456 (4)	0.01/1(2)	2.2(1)
Data collection		C(12)	0.3/28 (4)	0.2822 (4)	-0.0356 (2)	2.8(1)
Enrof Nonius CAD 4	$A = 70^{\circ}$	C(13)	0.4037 (4)	0.1750(5)	-0.0142(2)	3.1(1)
Elliai–ivolitus CAD-4	$\sigma_{\rm max} = 70$	C(14)	0.3010(4) 0.3247(4)	0.1247(4) 0.1867(4)	0.0387(2) 0.1102(2)	20(1)
diffractometer	$n = -10 \rightarrow 10$	C(16)	0.3247 (4)	0.1807(4)	0.0903(2)	2.5(1)
$\omega/2\theta$ scans	$k = -14 \rightarrow 14$	C(10)	0.1460 (3)	0.2902(4) 0.3745(4)	0.0903(2) 0.3942(2)	2.3(1) 20(1)
Absorption correction:	$l = -16 \rightarrow 16$	C(18)	0.2094(3)	0.6083(4)	0.4623(2)	2.3(1)
empirical	3 standard reflections	C(19)	0.1296 (3)	0.4119 (4)	0.5287 (2)	2.1 (1)
$T_{1} = 0.77 T_{1} = 1.00$	frequency: 60 min	C(20)	0.0446 (4)	0.1833 (4)	0.5585 (2)	2.4 (1)
$n_{\rm min} = 0.77$, $n_{\rm max} = 1.00$	interaction of the second second	C(21)	0.1107 (3)	0.3162 (4)	0.4657 (2)	2.1 (1)
9870 measured renections	intensity variation: -5.0%	C(22)	-0.5432 (3)	0.2724 (4)	0.4028 (2)	2.0(1)
5020 independent reflections	(linear correction	C(23)	-0.6175 (4)	0.0165 (4)	0.3695 (2)	2.5 (1)
4338 observed reflections	applied)	C(24)	-0.6414 (3)	0.0817 (4)	0.4880 (2)	2.2 (1)
$[I > 3\sigma(I)]$		C(25)	-0.6314 (4)	0.2083 (4)	0.5930 (2)	2.6(1)
$R_{i-1} = 0.067$		C(26)	-0.5857 (3)	0.2287 (4)	0.4762 (2)	2.1 (1)
ANDR 0.007						

0(9) O(10)

N(1)

Refinement

Refinement on F	Extinction correction:	Table	2. FI
R = 0.034	isotropic		isotr
wR = 0.048	(Zachariasen, 1963)		
S = 2.02	Extinction coefficient:		
4338 reflections	0.44323×10^{-5}		
431 parameters	Atomic scattering factors		
$w = 1/\sigma^2(F_o)$	from International Tables	S(1)	0.
$(\Delta/\sigma)_{\rm max} < 0.001$	for X-ray Crystallography	S(2)	0.
$\Delta \alpha = 0.48 \text{ e} ^{3}$	(1074 Vol IV)	O (1)	0.
$\Delta p_{\text{max}} = 0.48 \text{ C A}$	(17/4, 01.1)	O(2)	0.
$\Delta \rho_{\rm min} = -0.51 {\rm e A}^{\circ}$		O(3)	0.
		O(4)	0.
		O(5)	0.
Table 1. Fractional atom	ic coordinates and equivalent	O(6)	0.
• , • ,• ,		O(7)	0.
isotropic displacem	ent parameters (A ²) for	O(8W)	0.
•	~		

Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters $(Å^2)$ for Orange G/adenine

$B_{\rm eq} = (8\pi^2/3) \sum_i \sum_j U_{ij} a_i^* a_j^* \mathbf{a}_i \cdot \mathbf{a}_j.$

		on / o/_i_jo	-, -, -, -, -, -, -, -, -, -, -, -, -, -		N(2)
	x	у	Z	Beq	N(3)
S(1)	0.27564 (8)	0.63144 (9)	0.20474 (4)	1.97 (2)	N(4)
S(2)	-0.03967 (8)	0.9206(1)	0.23031 (4)	1.95 (2)	N(5)
O(1)	0.3035 (2)	0.7057 (3)	0.2822(1)	2.16 (7)	N(6)
O(2)	0.1767 (3)	0.4737 (3)	0.2054 (1)	2.94 (8)	N(7)
0(3)	0.4151 (3)	0.6699 (3)	0.1709(1)	2.56 (8)	N(8)
O(4)	-0.0866 (3)	1.0100 (3)	0.1862(1)	2.89 (9)	C(1)
O(5)	0.0804 (3)	1.0081 (3)	0.2897 (1)	2.59 (8)	C(2)
O(6)	-0.1628 (3)	0.7966 (3)	0.2612(1)	3.07 (8)	C(3)
O (7)	0.2449 (3)	0.5869 (3)	-0.1159 (1)	3.15 (9)	C(4)

ractional atomic coordinates and equivalent **m** 11 opic displacement parameters (Å²) for Orange G/cytosine

$B_{\rm eq} = (8\pi^2/3) \sum_i \sum_i U_{ij} a_i^* a_i^* \mathbf{a}_i \cdot \mathbf{a}_j.$

		-	
x	у	Ζ	Beq
0.53621 (4)	1.25003 (3)	0.74078 (2)	1.23(1)
0.22690 (4)	0.80763 (3)	0.89456 (2)	1.21 (1)
0.4539(1)	1.2890(1)	0.66224 (8)	1.81 (4)
0.7104(1)	1.3114 (1)	0.74289 (8)	1.77 (4)
0.4672(1)	1.2602(1)	0.83575 (8)	1.66 (4)
0.1761 (1)	0.6777 (1)	0.86761 (8)	1.68 (4)
0.0955(1)	0.8527 (1)	0.90436 (9)	1.89 (4)
0.3342 (1)	0.8382 (1)	0.98040 (8)	1.57 (4)
0.8522 (2)	1.0787 (1)	0.45398 (9)	2.09 (4)
0.5542 (2)	1.5524 (1)	0.7039 (1)	2.24 (5)
0.1285 (2)	1.2631 (1)	1.29381 (9)	2.07 (4)
0.4421 (1)	1.3182 (1)	1.07561 (9)	1.94 (4)
0.6892 (2)	1.2087 (1)	0.55561 (9)	1.28 (4)
0.7869 (2)	1.2658 (1)	0.4876(1)	1.43 (4)
0.3103 (2)	1.0500 (2)	1.0718(1)	1.88 (5)
0.2141 (2)	1.1542(1)	1.1838 (1)	1.42 (4)
0.0070 (2)	1.0538 (2)	1.2882(1)	1.77 (5)
0.8574 (2)	1.5127 (2)	0.8743 (1)	1.86 (5)
0.6475 (2)	1.4196 (1)	0.9766 (1)	1.37 (4)
0.6302 (2)	1.4968 (1)	1.1302(1)	1.65 (5)
0.6726 (2)	1.0922 (2)	0.5761 (1)	1.23 (5)
0.7487 (2)	1.0241 (2)	0.5177 (1)	1.49 (5)
0.7011 (2)	0.8939 (2)	0.5306(1)	1.61 (5)
0.5905 (2)	0.8358 (2)	0.5974 (1)	1.46 (5)

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C(5)	0.5197 (2)	0.9009 (2)	0.6610(1)	1.22 (5)
C(6)	0.4122 (2)	0.8313 (2)	0.7317(1)	1.32 (5)
C(7)	0.3476 (2)	0.8894 (2)	0.7973 (1)	1.27 (5)
C(8)	0.3893 (2)	1.0169 (2)	0.7938(1)	1.31 (5)
C(9)	0.4938 (2)	1.0879 (1)	0.7249(1)	1.12 (5)
C(10)	0.5622 (2)	1.0305 (2)	0.6537(1)	1.14 (5)
C(11)	0.8004 (2)	1.3847 (2)	0.4584(1)	1.42 (5)
C(12)	0.7114 (2)	1.4469 (2)	0.5018(1)	1.79 (5)
C(13)	0.7323 (2)	1.5642 (2)	0.4717(1)	2.10 (6)
C(14)	0.8420 (2)	1.6200 (2)	0.3991 (1)	2.14 (6)
C(15)	0.9287 (2)	1.5566 (2)	0.3554(1)	2.01 (6)
C(16)	0.9087 (2)	1.4380 (2)	0.3842(1)	1.72 (5)
C(17)	0.2074 (2)	1.0477 (2)	1.1403 (1)	1.51 (5)
C(18)	0.0892 (2)	0.9365 (2)	1.1731 (1)	2.30 (6)
C(19)	-0.0070 (2)	0.9454 (2)	1.2460(1)	2.23 (6)
C(20)	0.1150 (2)	1.1639 (2)	1.2576(1)	1.54 (5)
C(21)	0.7876 (2)	1.5120 (2)	0.9582(1)	1.50 (5)
C(22)	0.8502 (2)	1.6053 (2)	1.0307(1)	1.87 (5)
C(23)	0.7674 (2)	1.5942 (2)	1.1143 (1)	1.98 (6)
C(24)	0.5668 (2)	1.4059 (2)	1.0631 (1)	1.45 (5)

Table 3. Selected geometric parameters (Å, °) for

C(23)	0.7074(2)	1.3942	(2)	1.1145(1)	1.98(0)			0		
C(24)	0.5668 (2)	1.4059	(2)	1.0631 (1)	1.45 (5)	S(1)O(1)		1.456(1)	N(8)—C(24)	1.355 (2)
						S(1)O(2)		1.458(1)	C(1)—C(2)	1.456 (2)
Table	2 Salaat	ad a company		ana at ana (Å	a) fam	S(1)O(3)		1.468(1)	C(1)-C(10)	1.463 (2)
Table .	5. Select	ea geomen	ne pare	imeters (A,	°)jor	S(1)—C(9)		1.798 (2)	C(2)—C(3)	1.431 (2)
		Orange G	ladenir	ne		S(2)O(4)		1.459(1)	C(3)—C(4)	1.347 (2)
				- 		S(2)O(5)		1.451(1)	C(4) - C(5)	1.443 (2)
S(1) = O(1)		1.471 (2)	N(11)	C(24)	1.369 (4)	S(2)O(6)		1.466 (1)	C(5)—C(6)	1.409 (2)
S(1) = O(2)		1.453 (3)	N(11)	C(25)	1.352 (5)	S(2)-C(7)		1.775(2)	C(5) - C(10)	1.421 (2)
S(1) = O(3)		1.450 (3)	N(12)—	C(25)	1.330 (5)	O(7) - C(2)		1.273 (2)	$C(6) \rightarrow C(7)$	1.372 (2)
S(1) - C(9)		1.792 (4)	N(12)—	C(26)	1.384 (5)	O(9) - C(20)		1.219(2)	$C(7) \rightarrow C(8)$	1 396 (2)
S(2)O(4)		1.454 (4)	C(1)—C	(2)	1.455 (4)	O(10) - C(24)		1224(2)	$C(8) \rightarrow C(9)$	1.390(2)
S(2)O(5)		1.457 (2)	C(1)—C	(10)	1.462 (5)	N(1) - N(2)		1 295 (2)	C(0) = C(10)	1 4 3 9 (2)
S(2)O(6)		1.458 (3)	C(2)—C	(3)	1.428 (7)	$N(1) \rightarrow C(1)$		1.295(2)	C(1) - C(12)	1 387 (2)
S(2)—C(7)		1.763 (4)	C(3)—C	(4)	1.344 (6)	N(2) = C(11)		1.941(2) 1.404(2)	C(11) = C(12)	1.367(2)
O(7)—C(2)		1.279 (5)	C(4)—C	(5)	1.450 (4)	N(2) = C(17)		1 313 (2)	C(12) = C(13)	1.377 (2)
N(1)—N(2)		1.298 (5)	C(5)—C	(6)	1.391 (5)	N(3) = C(17)		1.313(2)	C(12) = C(13)	1.373 (3)
N(1) - C(1)		1.332 (6)	C(5)—C	(10)	1.416 (6)	N(4) = C(17)		1.333(2)	C(13) = C(14)	1.395 (2)
N(2)—C(11)		1.395 (6)	C(6)—C	(7)	1.382 (4)	N(4) = C(20)		1.361 (2)	$C(14) \rightarrow C(13)$	1.385 (3)
N(3)-C(17)		1.309 (5)	C(7)—C	(8)	1.390 (6)	N(5) = C(19)		1.330(3)	C(13) = C(16)	1.385 (3)
N(4)—C(17)		1.367 (5)	C(8)-C	(9)	1.382 (5)	N(3) = C(20)		1.308(2)	$C(17) \rightarrow C(18)$	1.423 (3)
N(4)-C(18)		1.350 (5)	C(9)-C	(10)	1.438 (4)	N(0) - C(21)		1.312(2)	C(18) - C(19)	1.352 (2)
N(5)-C(18)		1.320 (4)	C(11)(C(12)	1.398 (6)	N(7) = C(21)		1.351 (2)	C(21) - C(22)	1.423 (2)
N(5)-C(19)		1.350 (5)	cin	C(16)	1.397 (5)	N(7) - C(24)		1.3/4(2)	C(22) - C(23)	1.351 (2)
N(6)-C(19)		1.366 (4)	C(12)-	C(13)	1.375 (7)	N(8) - C(23)		1.358(2)		
N(6)-C(20)		1.372 (5)	C(13)	C(14)	1.387 (6)	O(1) - S(1) - O(1)	D(2)	114.65 (8)	C(6)—C(7)—C(8)	119.9 (1)
N(7) - C(20)		1.316 (4)	C(14)	C(15)	1.380(7)	O(1)-S(1)O	D(3)	110.87 (7)	C(7) - C(8) - C(9)	121.9(1)
N(7) - C(21)		1.380 (5)	C(15)	C(16)	1.377 (7)	O(1) - S(1) - O(1) -	C(9)	107.57 (8)	S(1) - C(9) - C(8)	113.0(1)
N(8) - C(22)		1.314 (5)	$C(17) \rightarrow 0$	C(21)	1.408 (4)	O(2)-S(1)-O	D (3)	111.06 (7)	S(1) - C(9) - C(10)	126.9 (1)
N(9) - C(22)		1.358 (5)	C(19)-	C(21)	1 386 (5)	O(2) - S(1) - O(2) - O(2) - S(1) - O(2) -	2(9)	107.56 (7)	C(8) - C(9) - C(10)	120.1 (1)
N(9) - C(23)		1.356 (5)	C(22)	C(26)	1 409 (4)	O(3) - S(1) - O(3) -	2(9)	104.50(7)	C(1) - C(10) - C(5)	117.0(1)
N(10) - C(23)		1 315 (4)	C(24)	C(26)	1 373 (5)	O(4) - S(2) - O(4) - S(2) - O(4) - S(2) - O(4) - S(2) - O(4) -	O(5)	114.08 (8)	C(1) - C(10) - C(9)	1267(1)
N(10) - C(24)		1 365 (5)	0(21)	C(20)	1.575 (5)	O(4) - S(2) - O(4) - S(2) - O(4) - S(2) - O(4) - S(2) - O(4) -	D(6)	112.33 (7)	C(5) - C(10) - C(9)	1164(1)
						O(4) - S(2) - O(4) - S(2) - O(4) - S(2) - O(4) - S(2) - O(4) -	C(7)	105.78 (8)	N(2) - C(11) - C(12)	121.6(1)
O(1) - S(1) - O(1) -	J(2)	110.5 (1)	C(7)—C	(8)—C(9)	122.2 (3)	O(5) - S(2) - O(5) -	D(6)	111 85 (7)	N(2) = C(11) = C(12)	1169(1)
O(1) - S(1) - O(1) -	J(3)	111.0(1)	S(1) - C((9)—C(8)	112.9 (2)	O(5) - S(2) - C	(0)	107 23 (7)	$C(12) \rightarrow C(11) \rightarrow C(16)$	121 4 (2)
O(1) - S(1) - O(1) -	C(9)	104.5 (2)	S(1) - C((9)—C(10)	127.3 (3)	O(6) - S(2) - C	(7)	104 80 (7)	C(11) - C(12) - C(13)	121.4(2)
O(2) - S(1) - O(2) - S(1) - O(2) - S(1) - O(2) - S(1) - O(2) -	D(3)	114.9 (2)	C(8)—C	(9)—C(10)	119.9 (3)	N(2) - N(1) - 0	Cú	119.0(1)	C(12) - C(13) - C(14)	1204(2)
O(2) = S(1) = O(2)	2(9)	106.8 (2)	C(1)—C	(10) - C(5)	117.9 (3)	N(1) - N(2) - 0		1214(1)	C(13) = C(14) = C(15)	120.4(2)
O(3) - S(1) - O(3) - S(1) - O(3) - S(1) - O(3) - S(1) - O(3) -	2(9)	108.5 (2)	C(1)—C	(10)—C(9)	125.8 (4)	C(17) = N(4)	-C(20)	125 3 (2)	C(14) - C(15) - C(16)	120.1(2) 120.4(2)
O(4) - S(2) - O(4) - S(2) - O(4) - S(2) - O(4) - S(2) - O(4) -	J(5)	112.8 (2)	C(5)—C	(10)—C(9)	116.3 (3)	C(19) - N(5)	-C(20)	122.5(2)	C(11) - C(16) - C(15)	120.4(2)
O(4)S(2)(D(6)	113.7 (2)	N(2)—C	(11)—C(12)	117.6 (3)	C(21) = N(7)	-C(24)	122.0(1)	N(3) = C(17) = N(4)	110.0(2)
O(4)S(2)C	C(7)	106.6 (2)	N(2)—C	(11)—C(16)	122.0 (4)	C(23) = N(8)	-C(24)	127.3(2)	N(3) = C(17) = C(18)	119.7(2)
O(5)S(2)(D(6)	111.3 (1)	C(12)(C(11)—C(16)	120.3 (4)	N(1) = C(1) = C(1)	2(2+)	122.3(1)	N(4) = C(17) = C(18)	122.0(2)
O(5)—S(2)—C	C(7)	105.8 (2)	C(11)—(C(12)C(13)	119.4 (3)	N(1) = C(1) = 0	2(2)	121.0(1)	C(17) = C(18) = C(10)	117.7(1)
O(6)S(2)(C(7)	106.0 (2)	C(12)—(C(13)—C(14)	120.5 (4)	C(2) = C(1) = C(1)	2(10)	120.2 (1)	V(1) - V(10) - V(19)	117.5(2)
N(2)—N(1)—4	C(1)	119.0 (3)	C(13)(C(14)—C(15)	119.8 (5)	C(2) = C(1) = C(2)	2(10)	120.2(1)	O(0) = O(19) = O(18)	122.0(2)
N(1)N(2)	C(11)	121.0 (3)	C(14)	C(15)—C(16)	121.0 (4)	0(7) - 0(2) -	(1)	120.0(2)	O(9) = O(20) = N(4)	121.7(2)
C(17)—N(4)—	-C(18)	124.8 (3)	C(11)—C	C(16)—C(15)	119.0 (4)	C(1) = C(2) = C(2)	(3)	120.4(1)	N(4) = C(20) = N(5)	125.6(1)
C(18)—N(5)—	-C(19)	111.8 (3)	N(3)—C	(17)—N(4)	121.2 (3)	$C(1) \rightarrow C(2) \rightarrow C(2)$	(3)	119.0(1)	N(4) = C(20) = N(3)	114.5 (2)
C(19)—N(6)—	-C(20)	106.4 (3)	N(3)—C	(17)—C(21)	126.1 (4)	C(2) - C(3) -	-(4) 	120.5(1)	N(0) - C(21) - N(7)	118.9 (2)
C(20)—N(7)—	-C(21)	102.8 (3)	N(4)—C	(17)—C(21)	112.7 (3)	C(3) = C(4) = C(4)	2(3)	122.3 (2)	N(0) = C(21) = C(22)	123.3 (2)
C(22)—N(9)—	-C(23)	124.4 (3)	N(4)—C	(18)—N(5)	124.8 (3)	$C(4) \rightarrow C(5) \rightarrow C(5)$	2(0) 2(10)	117.2(1)	N(7) = C(21) = C(22)	117.8(1)
C(23)—N(10)-	C(24)	111.4 (3)	N(5)—C	(19)—N(6)	127.6 (3)	C(4) - C(3) - C(3)	2(10)	120.0(1)	C(21) - C(22) - C(23)	117.8(2)
C(24)-N(11)-	-C(25)	105.9 (3)	N(5)-C	(19)—C(21)	127.6 (3)		J(10)	122.1(1)	N(8) - C(23) - C(22)	121.8 (2)
C(25)-N(12)-	-C(26)	102.9 (3)	N(6)—C	(19)—C(21)	104.9 (3)		-(/) 	119.7 (2)	O(10) - C(24) - N(7)	120.8 (2)
N(1)C(1)C	C(2)	122.6 (3)	N(6)—C	(20)—N(7)	113.9 (3)	S(2) - C(7) - C	.(D) 	121.1(1)	U(10) - U(24) - N(8)	123.2(1)
N(1)-C(1)-C	C(10)	117.0 (3)	N(7)—C	(21)—C(17)	129.6 (3)	S(2)-C(/)-C	.(0)	118.7(1)	IN(7)—C(24)—N(8)	116.0 (2)
C(2)—C(1)—C	C(10)	120.2 (4)	N(7)—C	(21)—C(19)	112.0 (3)		N(1)-N(2)C(11)C(1	2) 0.7 (2))
O(7)-C(2)-C	C(1)	121.0 (4)	C(17)-C	C(21)—C(19)	118.4 (3)		N(2)—N(1)—C(1)—C(2)	-6.3 (2)	1
O(7)C(2)C	C(3)	120.5 (3)	N(8)-C	(22)—N(9)	120.5 (3)		C(1)-N(1)	-N(2)-C(11) 175.5 (1)	

C(1)—C(2)—C(3)	118.4 (4)	N(8)-C(22)-C(26)	126.1 (3)
C(2)—C(3)—C(4)	120.7 (3)	N(9)-C(22)-C(26)	113.4 (3)
C(3)—C(4)—C(5)	123.0 (4)	N(9)-C(23)-N(10)	124.9 (3)
C(4)—C(5)—C(6)	118.7 (4)	N(10)-C(24)-N(11)	126.0 (3)
C(4)—C(5)—C(10)	119.1 (3)	N(10)-C(24)-C(26)	127.8 (3)
C(6)—C(5)—C(10)	122.2 (3)	N(11)—C(24)—C(26)	106.2 (3)
C(5)—C(6)—C(7)	120.2 (4)	N(11)-C(25)-N(12)	114.0 (3)
S(2)—C(7)—C(6)	122.9 (3)	N(12)-C(26)-C(22)	131.1 (3)
S(2)-C(7)-C(8)	118.1 (2)	N(12)-C(26)-C(24)	110.9 (3)
C(6)—C(7)—C(8)	119.0 (3)	C(22)-C(26)-C(24)	118.0 (3)
N(1)N	I(2)—C(11)—C	C(12) - 177.8(2)	
N(2)N	I(1)—C(1)—C(2) 7.5 (4)	
C(1)N	I(1)—N(2)—C(11) -177.1 (2)	

Table 4. Selected geometric parameters (Å, °) for Orange G/cytosine

Table 5. Close intermolecular contacts (Å)

Orange G/Adenine		Orange G/Cytosine			
O(1)…N(4)	2.742 (4)	O(1)…O(8W)	2.920 (2)		
$D(2) \cdots N(3)$	2.809 (4)	O(2)…N(6)	2.857 (2)		
$O(3) \cdots O(12W)$	2.850 (5)	O(3)N(7)	2,740 (2)		
O(3)O(10W)	2.988 (5)	O(4)N(6 ^{iv})	2.803 (2)		
O(4)…O(12W)	2.778 (5)	O(5)…N(4 ^{vii})	2.971 (2)		
O(5)…N(3")	2.824 (5)	O(6)…N(3)	2.837 (2)		
O(6)O(10W)	2.774 (4)	O(7)…N(5 ^{viii})	2.740 (2)		
O(7)O(11W ^{/iii})	2.928 (4)	O(8W)N(8ix)	2.761 (2)		
O(8W)N(6)	2.873 (4)	O(8W)O(9ix)	2.873 (2)		
O(9W)N(11)	2.760 (5)	O(10)N(4)	2.699 (2)		
O(9W)…N(5 ^{iv})	2.878 (5)	O(10)…N(3)	2.915 (3)		
O(10W)N(8)	2.855 (4)	.,			
$O(11W) \cdots O(12W)$	2.635 (7)				
O(11W)…N(9)	2.694 (4)				
N(8)…N(12 ^{vi})	2.882 (5)				

Symmetry codes: (i) 1 + x, y, z; (ii) x, 1 + y, z; (iii) -x, 1 - y, -z; (iv) -1 + x, -1 + y, z; (v) x, -1 + y, z; (vi) -1 - x, 1 - y, 1 - z; (vii) -x, 2 - y, 2 - z; (viii) 1 + x, y, -1 + z; (ix) 1 - x, 3 - y, 2 - z.

In both structures, H atoms bonded to C atoms were placed in calculated positions and were not refined. H atoms bonded to N atoms in Orange G/adenine were located in difference maps and their positional parameters were refined. Peaks were also found near the water O atoms, but their positions were not considered chemically reasonable and were left unassigned. H atoms bonded to N atoms and to the water O atom in Orange G/cytosine were located and their positional parameters were refined. Refinement of a secondary extinction coefficient (Zachariasen, 1963) was considered warranted only in the case of Orange G/cytosine. Data collection and cell determination were carried out using Enraf-Nonius CAD-4 software. Data reduction and structure refinement used the TEXSAN software package (Molecular Structure Corporation, 1985). Structure solution was accomplished using SHELXS86 (Sheldrick, 1985). Molecular graphics were prepared using ORTEPII (Johnson, 1976) and PLUTO (Motherwell & Clegg, 1978).

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: BK1002). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Sorbic Acid

Philip J. Cox

School of Pharmacy, The Robert Gordon University, Schoolhill, Aberdeen AB9 1FR, Scotland

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Abstract

The crystal and molecular structure of sorbic acid [(E,E)-2,4-hexadienoic acid], $C_6H_8O_2$, has been determined. There appears to be no delocalization of the C_{sp^2} — C_{sp^2} bonds, arranged in an *E,E* configuration, which have lengths of 1.328 (2) and 1.335 (2) Å (double) and 1.442 (2)–1.487 (2) Å (single). Dimers are formed in the crystal with an intermolecular O…O separation of 2.632 (2) Å; O—H 0.99 (2), H…O 1.65 (2) Å, O—H…O 174 (2)°.

Comment

Sorbic acid, (I), exhibits antibacterial and antifungal properties (Martindale, 1993) and has been used to prevent spoilage of syrup by moulds (Richards, 1972). The structures of numerous simple carboxylic acids have been determined and many involve intermolecular hydrogen bonding (Speakman, 1972; CSSR, 1993). It is difficult to obtain well formed crystals of sorbic acid and attempts to determine the cell dimensions by X-ray powder diffraction from a commercial crystalline sample (99 + %, Sigma) were unsuccessful. After repeated crystallizations from ethanol only one specimen was considered suitable for data collection.



The study shows that dimers form in the crystal with intermolecular hydrogen bonding around a centre of symmetry. The short O…O separation is