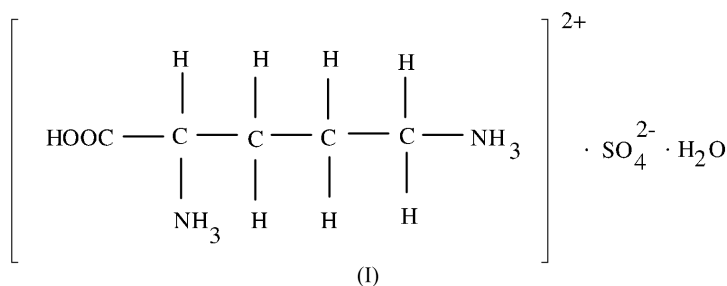


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rkrspmku@yahoo.co.in**Key indicators**Single-crystal X-ray study  
 $T = 293$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.008$  Å  
 $R$  factor = 0.059  
 $wR$  factor = 0.171  
Data-to-parameter ratio = 8.6For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.**L-Ornithinium sulfate monohydrate**

In the title compound,  $\text{C}_5\text{H}_{14}\text{N}_2\text{O}_2^{2+} \cdot \text{SO}_4^{2-} \cdot \text{H}_2\text{O}$ , the ornithinium cation forms a strong  $\text{O}-\text{H} \cdots \text{O}$  hydrogen bond with the sulfate anion. One backbone conformation angle is in an eclipsed conformation, while the other two are in a *trans-trans* staggered conformation. All the amino and carboxylate groups and the sulfate anions are involved in hydrogen bonding. The amino groups are also involved in a three-centred hydrogen bond with the O atoms of the sulfate anion.

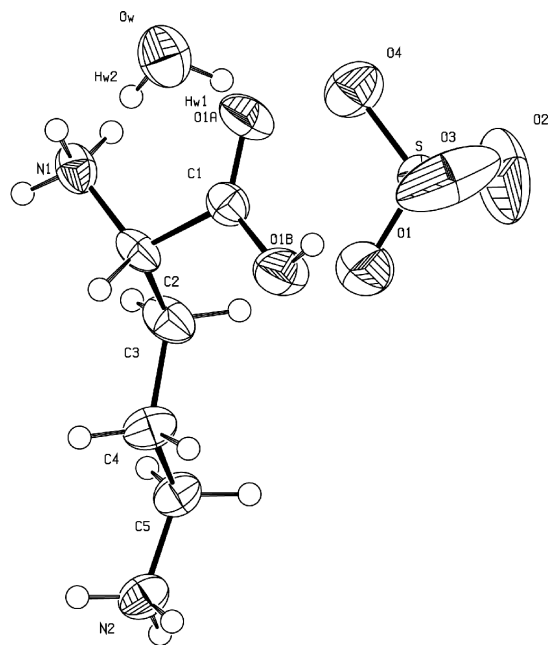
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Online 22 October 2004**Comment**

L-Ornithine helps to elevate levels of human growth hormone, assists in repairing muscles and tissues, stimulates the pancreas to release insulin and helps regenerate the liver, in addition to being necessary for the proper functioning of the immune system (Bucchi *et al.*, 1990). The crystal structures of L-ornithine hydrochloride (Chiba *et al.*, 1967), L-ornithine nitrate (Ramaswamy *et al.*, 2002) and bis(L-ornithinium) chloride nitrate sulfate (Ramaswamy *et al.*, 2004) have been solved. In the present study, the crystal structure of L-ornithinium sulfate monohydrate, (I) (Fig. 1), obtained by the recrystallization of L-ornithine from sulfuric acid, is reported.

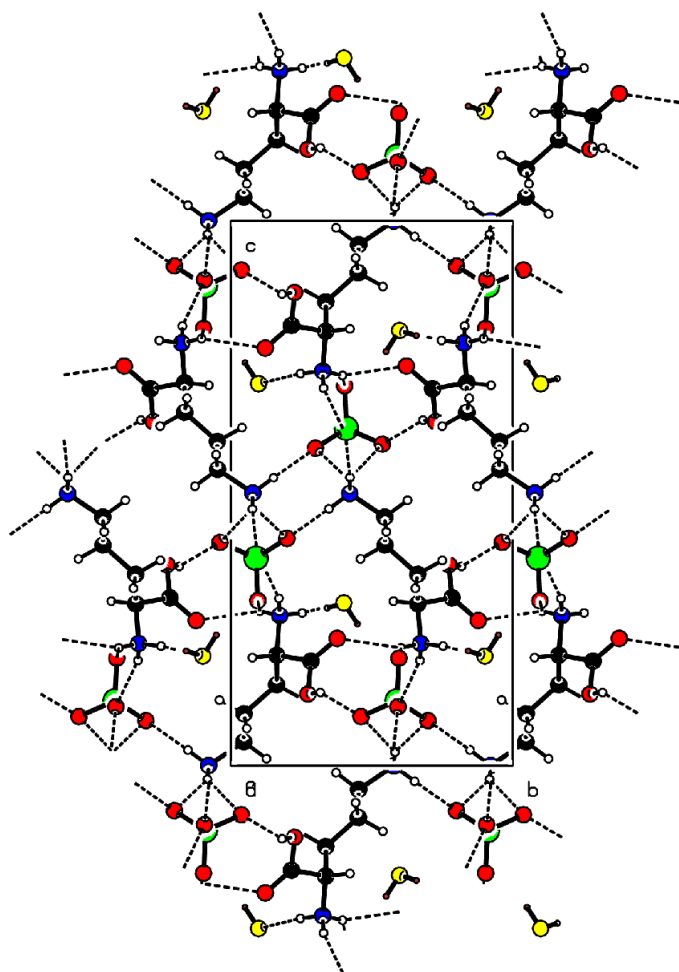


The average S—O bond distance and O—S—O bond angle in (I) conform to tetrahedral symmetry. The unsymmetrical C—O bond distances [1.1957 (6) and 1.3162 (7) Å] and O—C—C bond angles [123.0 (5) and 111.5 (5)°] clearly confirm the protonation of the carboxyl group. Both amino groups are also protonated.

The backbone conformation angle  $\psi^1$  (O1A—C1—C2—N1) indicates a *cis* conformation [−9.3 (8) Å]. The deviation of this amino N atom from the carbonyl plane is very small [0.208 (9) Å]. An eclipsed conformation is observed about C2—C3, with torsion angles N1—C2—C3—C4 = −130.3 (5)° and C1—C2—C3—C4 = 110.4 (6)°. The other angles,  $\chi^2$  (C2—C3—C4—C5) and  $\chi^3$  (C3—C4—C5—N2), have fully extended *trans-trans* conformations [178.2 (5) and −170.8 (5)°, respectively]. The maximum deviation of the side-chain atoms from their mean plane is 0.41 (1) Å for C3.



**Figure 1**  
The molecular structure of (I), with the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.



**Figure 2**  
A packing diagram for (I), viewed down the *a* axis. Dashed lines indicate the hydrogen bonds.

The hydroxyl group of the ornithinium cation forms a strong O—H···O hydrogen bond with one O atom of the sulfate anion (Fig. 2, Table 2). Atoms N1 and N2 form one two-centred and two three-centred hydrogen bonds, leading to class III behaviour (Jeffrey & Saenger, 1991). Both N atoms in the ornithinium cation connect O atoms of two different sulfate anions, thus forming a chain running along the *a* axis (N1—H1A···O4<sup>iii</sup> and N1—H1C···O3<sup>ii</sup>; Table 2) and a similar zigzag head-to-tail sequence along the *a* axis (N2—H2A···O3<sup>iv</sup> and N2—H2C···O2<sup>vi</sup>; Table 2). The water molecule connects two ornithinium cations to form an infinite chain, also along the *a* axis (N1—H1B···OW—HW2···O1A<sup>vii</sup>; Table 2). Through the three-centred bond, this water molecule connects two  $2_1$  screw-related sulfate anions along the *b* axis.

## Experimental

The title compound was formed by mixing a 1 M solution of L-ornithine with a 1 M solution of sulfuric acid, followed by crystallization by slow evaporation under ambient conditions.

### Crystal data

$C_5H_{14}N_2O_2^{2+} \cdot SO_4^{2-} \cdot H_2O$   
 $M_r = 248.26$   
 Orthorhombic,  $P2_12_12_1$   
 $a = 6.1589$  (2) Å  
 $b = 9.682$  (6) Å  
 $c = 18.714$  (6) Å  
 $V = 1115.9$  (8) Å<sup>3</sup>  
 $Z = 4$   
 $D_x = 1.478$  Mg m<sup>-3</sup>  
 $D_m = 1.475$  (5) Mg m<sup>-3</sup>

$D_m$  measured by flotation, using a mixture of CCl<sub>4</sub> and xylene  
 Mo  $K\alpha$  radiation  
 Cell parameters from 25 reflections  
 $\theta = 11.3$ – $14.2^\circ$   
 $\mu = 0.31$  mm<sup>-1</sup>  
 $T = 293$  (2) K  
 Block, colourless  
 $0.3 \times 0.3 \times 0.2$  mm

### Data collection

Nonius MACH 3 diffractometer  
 $\omega/2\theta$  scans  
 Absorption correction:  $\psi$  scan  
 (North *et al.*, 1968)  
 $T_{\min} = 0.911$ ,  $T_{\max} = 0.940$   
 1399 measured reflections  
 1172 independent reflections  
 1043 reflections with  $I > 2\sigma(I)$

$R_{\text{int}} = 0.029$   
 $\theta_{\max} = 25^\circ$   
 $h = 0 \rightarrow 7$   
 $k = -1 \rightarrow 11$   
 $l = -1 \rightarrow 22$   
 3 standard reflections  
 frequency: 60 min  
 intensity decay: none

### Refinement

Refinement on  $F^2$   
 $R[F^2 > 2\sigma(F^2)] = 0.059$   
 $wR(F^2) = 0.171$   
 $S = 1.06$   
 1172 reflections  
 137 parameters  
 H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.112P)^2 + 1.3829P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{\max} = 0.005$   
 $\Delta\rho_{\max} = 0.73$  e Å<sup>-3</sup>  
 $\Delta\rho_{\min} = -0.37$  e Å<sup>-3</sup>

**Table 1**

Selected geometric parameters (Å, °).

O1B—C1	1.319 (7)	O1A—C1	1.197 (7)
O1A—C1—O1B	125.3 (6)	O1B—C1—C2	111.6 (5)
O1A—C1—C2	123.1 (5)		
O1A—C1—C2—N1	−9.3 (8)	C1—C2—C3—C4	110.4 (6)
O1B—C1—C2—N1	171.8 (5)	C2—C3—C4—C5	178.2 (5)
N1—C2—C3—C4	−130.3 (5)	C3—C4—C5—N2	−170.8 (5)

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

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$O1B-H1\cdots O1^i$	0.82	1.87	2.546 (7)	139
$N1-H1A\cdots O1A^{ii}$	0.89	2.21	2.903 (6)	134
$N1-H1A\cdots O4^{iii}$	0.89	2.34	3.003 (7)	132
$N1-H1B\cdots OW$	0.89	1.91	2.790 (7)	169
$N1-H1C\cdots O3^{iv}$	0.89	1.99	2.810 (7)	153
$N2-H2A\cdots O3^{iv}$	0.89	1.86	2.729 (7)	165
$N2-H2B\cdots O2^v$	0.89	1.94	2.781 (8)	158
$N2-H2C\cdots O2^{vi}$	0.89	2.14	2.945 (10)	150
$N2-H2C\cdots O1^{vi}$	0.89	2.35	3.136 (8)	148
$OW-HW1\cdots O4$	0.88	1.88	2.726 (7)	162
$OW-HW2\cdots O4^{iii}$	0.87	2.46	3.195 (7)	143
$OW-HW2\cdots O1A^{vii}$	0.87	2.57	3.191 (7)	130

Symmetry codes: (i)  $x-1, y, z$ ; (ii)  $1-x, y-\frac{1}{2}, \frac{1}{2}-z$ ; (iii)  $2-x, y-\frac{1}{2}, \frac{1}{2}-z$ ; (iv)  $\frac{1}{2}+x, \frac{3}{2}-y, 1-z$ ; (v)  $x, y-1, z$ ; (vi)  $x-\frac{1}{2}, \frac{3}{2}-y, 1-z$ ; (vii)  $1+x, y, z$ .

For the water molecule, the H atoms were located in a difference Fourier map and in these positions. All remaining H atoms were positioned geometrically and refined in the riding-model approximation, with  $C-H = 0.97-0.98$ ,  $N-H = 0.89$  and  $O-H = 0.82$  Å, and with  $U_{iso}(H) = 1.2U_{eq}$  of the carrier atom ( $1.5U_{eq}$  for methyl and  $NH_3$  H atoms).

Data collection: *CAD-4 EXPRESS* (Enraf-Nonius, 1994); cell refinement: *CAD-4 EXPRESS*; data reduction: *XCAD4* (Harms & Wocadlo, 1995); program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEP3* (Farrugia, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97*.

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