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Acta Cryst. (1998). **C54**, 637–641

Complexes Between L-Leucine and its Precipitants

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(Received 16 December 1996; accepted 27 November 1997)

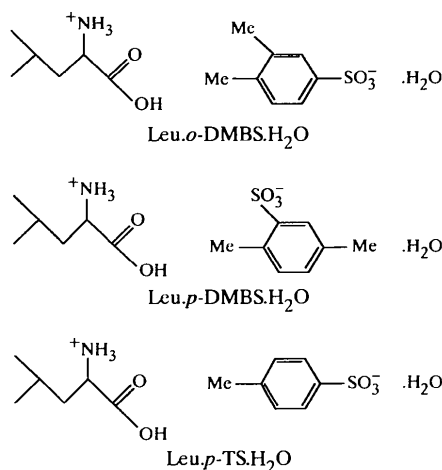
Abstract

The complexes between L-leucine and aromatic sulfonic acids, L-leucine 3,4-dimethylbenzenesulfonate monohydrate ($C_6H_{14}NO_2^+ \cdot C_8H_9O_3S^- \cdot H_2O$), L-leucine 2,5-dimethylbenzenesulfonate monohydrate ($C_6H_{14}NO_2^+ \cdot C_8H_9O_3S^- \cdot H_2O$), L-leucine 4-methylbenzenesulfonate monohydrate ($C_6H_{14}NO_2^+ \cdot C_7H_7O_3S^- \cdot H_2O$), as well as L-leucine benzenesulfonate monohydrate, exhibit a striking similarity in their packing schemes, which are dominated by van der Waals stacking of the hydrogen-bonded molecular double layers. The tightness of packing of double layers is most likely related to the solubility of the complexes.

Comment

L-Leucine (Leu) is one of the essential mammalian amino acids and is used for intravenous feeding solution.

Purification of Leu from hydrolysate of proteins is quite difficult because the hydrolysate contains a large amount of L-isoleucine (Ile) and L-valine (Val). These three amino acids have very similar crystal structures (Harding & Howieson, 1976; Torii & Iitaka, 1970, 1971), thus repetition of the recrystallization of Leu has been ineffective for purification because of formation of mixed crystals. Since the 1950s, many complexes between Leu and aromatic sulfonic acids (called precipitants) have been studied in order to purify Leu efficiently, and some of them are effective for the isolation of Leu. Among these precipitants, 3,4-dimethylbenzenesulfonic acid (Nagai, 1963) (*o*-DMBS) has extreme selectivity for Leu and has been used industrially. A number of precipitants resembling *o*-DMBS are known, *i.e.* 2,5-dimethylbenzenesulfonic acid (*p*-DMBS), 2,4-dimethylbenzenesulfonic acid (*m*-DMBS), *p*-toluenesulfonic acid (*p*-TS) (Hongo *et al.*, 1979) and benzenesulfonic acid (BS) (Hongo *et al.*, 1979). These precipitants form complexes with Leu, but the solubilities of the complexes are different to one another. The reduced solubilities based on Leu at 283 K are 0.82 g/100 g H₂O for the *o*-DMBS complex, 0.96 g/100 g H₂O for the *p*-DMBS complex, 6.24 g/100 g H₂O for the *m*-DMBS complex, 5.82 g/100 g H₂O for the *p*-TS complex and 7.66 g/100 g H₂O for the BS complex. The crystal structures of these five complexes were determined in order to elucidate the relationship between packing arrangement and functions as a precipitant, such as selectivity for Leu and reduced solubility. However, the results for the complex with BS have already been reported (Kimoto *et al.*, 1989) and the refinement of the complex with *m*-DMBS was unsatisfactory. Thus, we report here the crystal structures of the remaining three complexes and compare them with the results for the complex with BS.



All but one of these four complexes crystallize in the orthorhombic space group $P2_12_12_1$ with similar cell parameters. The crystal of the *m*-DMBS complex also

has the orthorhombic space group $P2_12_12_1$ with similar unit-cell parameters, *i.e.* $a = 9.839(2)$, $b = 27.86(1)$ and $c = 6.609(2)$ Å. Even though one of them has a different space group ($P2_1$), all the complexes consist of one Leu cation, one precipitant anion and one water molecule per asymmetric unit (Figs. 1, 2 and 3), and exhibit very similar packing schemes.

Hydrogen bonds hold the carboxyl and amino groups of Leu, the sulfonic acid group of the precipitants, and water together in corrugated double layers, while the hydrophobic side chain of Leu and the aromatic rings of the precipitants lie outside the layer (Figs. 4, 5 and 6). The crystal-packing arrangement consists of stacking of these double layers with van der Waals contacts.

All torsion angles of the side chain of Leu are similar to each other, and are also similar to those of free Leu, except for $\psi(O-C-C-N)$, *i.e.* -36 and -26° in the Leu crystal (Harding & Howieson, 1976). The $\psi(O-C-C-N)$ values of the complexes are $-6.4(6)$ and $173.1(4)^\circ$ for Leu.*o*-DMBS. H_2O , $-5.7(6)$ and $174.9(4)^\circ$ for Leu.*p*-DMBS. H_2O , and $-9.2(6)$ and $168.3(4)^\circ$ for Leu.*p*-TS. H_2O , so these differences seem to be caused by the hydrogen-bond scheme including the sulfonic acid group of the precipitants and water.

In the hydrophobic interaction area, some structural differences appear because of the different numbers and

positions of the methyl groups in the precipitants. In order to elucidate quantitatively the tightness of the packing of the complex molecule, we tried to calculate the filling factor in the unit cell using space-fill models and van der Waals radii for each atom (CRC Handbook

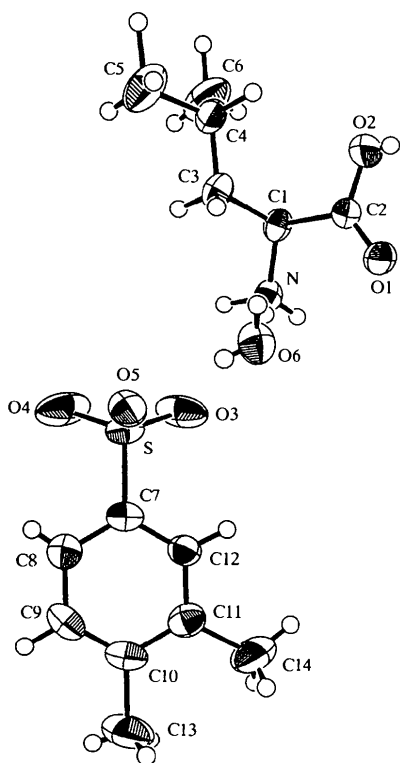


Fig. 1. ORTEP view (Johnson, 1976) of the Leu.*o*-DMBS. H_2O complex. Displacement ellipsoids are plotted at the 50% probability level.

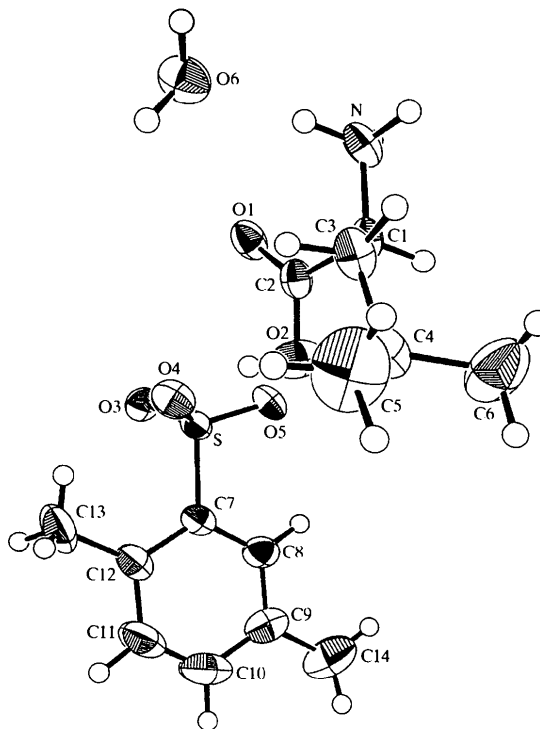


Fig. 2. ORTEP view (Johnson, 1976) of the Leu.*p*-DMBS. H_2O complex. Displacement ellipsoids are plotted at the 50% probability level.

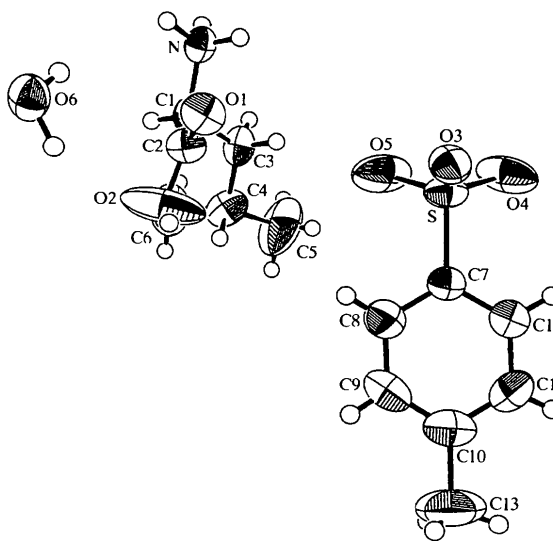


Fig. 3. ORTEP view (Johnson, 1976) of the Leu.*p*-TS. H_2O complex. Displacement ellipsoids are plotted at the 50% probability level.

of Chemistry and Physics, 1986). The results were 66.7% for the BS complex, 64.4% for the *p*-TS complex, 67.2% for the *p*-DMBS complex and 67.6% for the *o*-DMBS complex. These four complexes have similar packing of the hydrophilic areas, so the difference in the filling factors must be mainly due to the tightness of the packing of the hydrophobic areas. From these factors, it is seen that in the crystal of the complex with BS, there

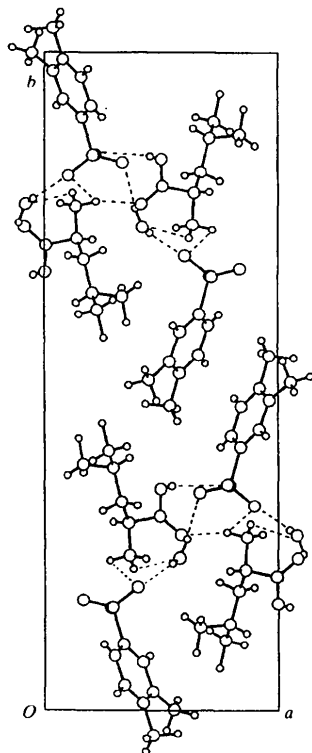


Fig. 4. Packing diagram of the *Leu.o*-DMBS.H₂O complex viewed down the *c* axis.

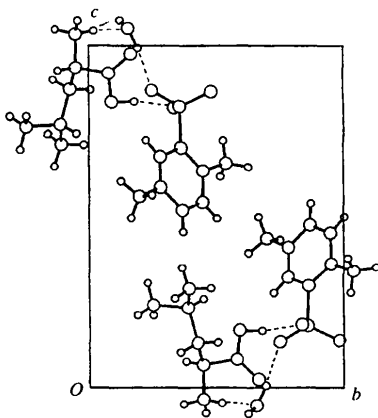


Fig. 5. Packing diagram of the *Leu.p*-DMBS.H₂O complex viewed down the *a* axis.

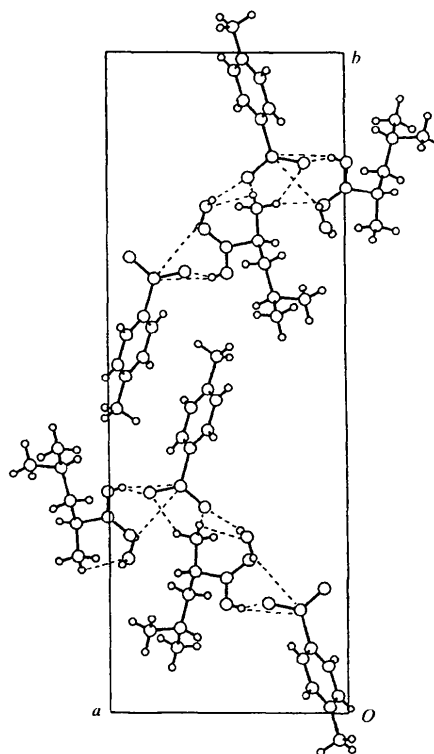


Fig. 6. Packing diagram of the *Leu.p*-TS.H₂O complex viewed down the *c* axis.

is a little cavity in the hydrophobic interaction area. In the case of the crystal of the complex with *p*-TS, having one more methyl group than the BS complex, the *b* axis is elongated by the addition of the methyl group, so that the cavity arises more in the hydrophobic area. In the crystals of the complexes with *o*- and *p*-DMBS, the size of the cavity is apparently decreased by the relatively close packing in these complexes. On the other hand, the filling factor of the *m*-DMBS complex remains 65.7%.

From a comparison of the filling factors of the complexes, it is reasonable that the complex of *Leu.o*-DMBS.H₂O has the lowest solubility in these complexes. It is suggested that the tightness of the packing of the hydrophobic region must play an important role in determining the solubility of the complexes.

Experimental

All precipitants were purchased and the crystals of the title complexes were obtained by slow cooling of equimolar warm aqueous solutions of L-leucine and the precipitants.

Leu.o-DMBS.H₂O

Crystal data

C₆H₁₄NO₂⁺·C₈H₉O₃S⁻·H₂O
M_r = 335.41

Cu Kα radiation
 λ = 1.5418 Å

Orthorhombic
*P*2₁2₁2₁
a = 9.855 (7) Å
b = 28.479 (9) Å
c = 6.280 (1) Å
V = 1763 (5) Å³
Z = 4
D_x = 1.264 Mg m⁻³
D_m = 1.268 Mg m⁻³
D_m measured by flotation
 in isooctane-carbon
 tetrachloride

Data collection

Rigaku AFC-5S diffractometer
 $\omega/2\theta$ scans
 Absorption correction:
 empirical via ψ scans
 (North *et al.*, 1968)
T_{min} = 0.485, *T_{max}* = 0.833
 2142 measured reflections
 2142 independent reflections

Refinement

Refinement on *F*
R = 0.044
wR = 0.044
S = 2.392
 1687 reflections
 199 parameters
 H atoms not refined
 Weighting scheme based
 on measured s.u.'s;
 $w = 1/\sigma^2(F)$

Cell parameters from 25
 reflections
 $\theta = 30\text{--}40^\circ$
 $\mu = 1.870 \text{ mm}^{-1}$
T = 298 K
 Plate
 0.7 × 0.5 × 0.1 mm
 Colourless

1687 reflections with
 $I > 3\sigma(I)$
 $\theta_{\text{max}} = 80^\circ$
 $h = 0 \rightarrow 13$
 $k = 0 \rightarrow 36$
 $l = 0 \rightarrow 8$
 3 standard reflections
 every 150 reflections
 intensity decay: none

$(\Delta/\sigma)_{\text{max}} = 0.052$
 $\Delta\rho_{\text{max}} = 0.27 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.23 \text{ e \AA}^{-3}$
 Extinction correction: none
 Scattering factors from *International Tables for X-ray Crystallography* (Vol. IV)

Monoclinic
*P*2₁
a = 6.163 (1) Å
b = 10.199 (9) Å
c = 14.140 (9) Å
 $\beta = 94.112 (9)^\circ$
V = 886.5 (1) Å³
Z = 2
D_x = 1.257 Mg m⁻³
D_m not measured

Data collection

Rigaku AFC-5S diffractometer
 $\omega/2\theta$ scans
 Absorption correction:
 empirical via ψ scans
 (North *et al.*, 1968)
T_{min} = 0.709, *T_{max}* = 0.834
 2142 measured reflections
 2142 independent reflections

Refinement

Refinement on *F*
R = 0.049
wR = 0.060
S = 3.729
 1803 reflections
 198 parameters
 H atoms not refined
 Weighting scheme based
 on measured s.u.'s;
 $w = 1/\sigma^2(F)$

Cell parameters from 25
 reflections
 $\theta = 30\text{--}40^\circ$
 $\mu = 1.860 \text{ mm}^{-1}$
T = 298 K
 Plate
 0.7 × 0.5 × 0.1 mm
 Colourless

1803 reflections with
 $I > 3\sigma(I)$
 $\theta_{\text{max}} = 80^\circ$
 $h = 0 \rightarrow 8$
 $k = 0 \rightarrow 13$
 $l = -18 \rightarrow 18$
 3 standard reflections
 every 150 reflections
 intensity decay: none

$(\Delta/\sigma)_{\text{max}} = 0.051$
 $\Delta\rho_{\text{max}} = 0.21 \text{ e \AA}^{-3}$
 $\Delta\rho_{\text{min}} = -0.50 \text{ e \AA}^{-3}$
 Extinction correction: none
 Scattering factors from *International Tables for X-ray Crystallography* (Vol. IV)

Table 2. Selected geometric parameters (Å, °) for *Leu.-p-DMBS.H₂O*

S—O3	1.458 (4)	C4—C5	1.535 (8)
S—O4	1.461 (3)	C4—C6	1.53 (1)
S—O5	1.461 (4)	C7—C8	1.391 (7)
S—C7	1.759 (4)	C7—C12	1.409 (7)
O1—C2	1.199 (6)	C8—C9	1.373 (7)
O2—C2	1.312 (6)	C9—C10	1.40 (1)
N—C1	1.478 (6)	C9—C14	1.49 (1)
C1—C2	1.518 (6)	C10—C11	1.35 (1)
C1—C3	1.546 (6)	C11—C12	1.387 (8)
C3—C4	1.519 (8)	C12—C13	1.504 (8)
O1—C2—C1—N	-5.7 (6)	N—C1—C3—C4	-164.3 (4)
O1—C2—C1—C3	114.5 (5)	C1—C3—C4—C5	-170.7 (5)
O2—C2—C1—N	174.9 (4)	C1—C3—C4—C6	66.9 (6)
O2—C2—C1—C3	-64.9 (6)	C2—C1—C3—C4	77.2 (6)

Leu.-p-TS.H₂O**Crystal data**

C₆H₁₄NO₂⁺.C₇H₇O₃S⁻.H₂O
M_r = 321.39
Orthorhombic
*P*2₁2₁2₁
a = 9.866 (2) Å
b = 28.201 (9) Å
c = 6.250 (2) Å
V = 1739 (1) Å³
Z = 4
D_x = 1.228 Mg m⁻³
D_m not measured

Cu *K*α radiation
 $\lambda = 1.5418 \text{ \AA}$
 Cell parameters from 25
 reflections
 $\theta = 30\text{--}40^\circ$
 $\mu = 1.875 \text{ mm}^{-1}$
T = 298 K
 Plate
 0.7 × 0.5 × 0.2 mm
 Colourless

Table 1. Selected geometric parameters (Å, °) for *Leu.-o-DMBS.H₂O*

S—O3	1.456 (4)	C4—C5	1.521 (7)
S—O4	1.455 (4)	C4—C6	1.518 (7)
S—O5	1.452 (3)	C7—C8	1.383 (7)
S—C7	1.768 (4)	C7—C12	1.375 (6)
O1—C2	1.195 (4)	C8—C9	1.387 (6)
O2—C2	1.296 (5)	C9—C10	1.378 (6)
N—C1	1.485 (5)	C10—C11	1.399 (7)
C1—C2	1.519 (5)	C10—C13	1.516 (6)
C1—C3	1.526 (6)	C11—C12	1.394 (5)
C3—C4	1.526 (6)	C11—C14	1.510 (7)
O1—C2—C1—N	-6.4 (6)	N—C1—C3—C4	-167.8 (3)
O1—C2—C1—C3	114.6 (5)	C1—C3—C4—C5	-178.3 (4)
O2—C2—C1—N	173.1 (4)	C1—C3—C4—C6	59.7 (5)
O2—C2—C1—C3	-65.9 (5)	C2—C1—C3—C4	73.6 (5)

Leu.-p-DMBS.H₂O**Crystal data**

C₆H₁₄NO₂⁺.C₈H₉O₃S⁻.H₂O
M_r = 335.41

Cu *K*α radiation
 $\lambda = 1.5418 \text{ \AA}$

Data collection

Rigaku AFC-5S diffractometer	1806 reflections with $I > 3\sigma(I)$
$\omega/2\theta$ scans	$\theta_{\max} = 80^\circ$
Absorption correction:	$h = 0 \rightarrow 13$
empirical via ψ scans	$k = 0 \rightarrow 36$
(North <i>et al.</i> , 1968)	$l = 0 \rightarrow 8$
$T_{\min} = 0.575$, $T_{\max} = 0.693$	3 standard reflections
2115 measured reflections	every 150 reflections
2115 independent reflections	intensity decay: none

Refinement

Refinement on F	$(\Delta/\sigma)_{\max} = 0.04$
$R = 0.047$	$\Delta\rho_{\max} = 0.23 \text{ e } \text{\AA}^{-3}$
$wR = 0.055$	$\Delta\rho_{\min} = -0.35 \text{ e } \text{\AA}^{-3}$
$S = 2.86$	Extinction correction: none
1806 reflections	Scattering factors from <i>International Tables for X-ray Crystallography</i> (Vol. IV)
189 parameters	
H atoms not refined	
Weighting scheme based on measured s.u.'s;	
$w = 1/\sigma^2(F)$	

Table 3. Selected geometric parameters (\AA , $^\circ$) for *Leu-p-TS.H₂O*

S—O3	1.447 (3)	C4—C5	1.512 (8)
S—O4	1.445 (3)	C4—C6	1.506 (7)
S—O5	1.442 (3)	C7—C8	1.370 (6)
S—C7	1.763 (4)	C7—C12	1.355 (6)
O1—C2	1.200 (4)	C8—C9	1.379 (6)
O2—C2	1.285 (5)	C9—C10	1.350 (7)
N—C1	1.472 (4)	C10—C11	1.367 (7)
C1—C2	1.515 (5)	C10—C13	1.516 (6)
C1—C3	1.529 (5)	C11—C12	1.386 (6)
C3—C4	1.504 (6)		
O1—C2—C1—N	-9.2 (6)	N—C1—C3—C4	-168.4 (3)
O1—C2—C1—C3	112.0 (5)	C1—C3—C4—C5	-176.0 (5)
O2—C2—C1—N	168.3 (4)	C1—C3—C4—C6	61.3 (6)
O2—C2—C1—C3	-70.5 (5)	C2—C1—C3—C4	72.6 (5)

Because of the poor ratio of the number of reflections to the number of parameters, the H-atom parameters were not included in the refinement.

For all compounds, data collection: *MSCIAFC Diffractometer Control Software* (Molecular Structure Corporation, 1988); cell refinement: *MSCIAFC Diffractometer Control Software*; data reduction: *MSCIAFC Diffractometer Control Software*; program(s) used to solve structures: *MITHRIL* (Gilmore, 1984); program(s) used to refine structures: *ALLS* (Lapp & Jacobson, 1979); software used to prepare material for publication: *PLUTO* (Motherwell & Clegg, 1983) and *ORTEPII* (Johnson, 1976).

The authors would like to thank Professor T. Yamane and Dr A. Suzuki of Nagoya University, Japan, for helpful advice. They also thank members of the Crystal Physics Laboratory of Tohoku University, Japan, and the editors of the IUCr.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: OAI024). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1998). **C54**, 641–643

10-Chloro-6,7-dihydro-5H-benzo[6,7]cyclohepta[1,2-b]quinoline and 10-Methyl-6,7-dihydro-5H-benzo[6,7]cyclohepta[1,2-b]quinoline

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(Received 5 June 1997; accepted 21 November 1997)

Abstract

The quinoline ring systems in the title molecules, C₁₈H₁₄ClN and C₁₉H₁₇N, are planar. The cycloheptane rings are in half-chair conformations and the crystal structures are stabilized by van der Waals interactions.

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