Metal—Penicillamine Interactions. Part 1. Preparation and Crystal Structure of a 1:1 Complex of Mercuric Chloride with D-Penicillamine, $2[(\mu_3-Cl){HgSC(CH_3)_2CH(NH_3)COO}_3] \cdot 3(\mu_2-Cl) \cdot 2(H_3O) \cdot (H_2O \cdot Cl)_3$

LANGFORD BOOK

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Received January 31, 1984

A 1:1 complex of mercuric chloride with Dpenicillamine has been isolated and characterised $2[(\mu_3-Cl){HgSC(CH_3)_2CH(NH_3)COO}_3] \cdot 3(\mu_2$ as Cl)·2(H₃O)·(H₂O·Cl)₃. The compound crystallises in cubic space group $P4_132$, with a = 18.679(5)Å and Z = 4. The structure, refined to $R_F = 0.086$ for 443 observed Mo-Ka diffractometer data, features a triply bridging chloride ion linking three equivalent $[HgSC(CH_3)_2 CH(NH_3)COO]^+$ units $[Hg-Cl = 2.37(1) \text{ Å}, Hg-Cl-Hg' = 98.5(9)^6]$. The carboxylate groups of a pair of adjacent penicillamine ligands are strongly linked via a symmetrical O···H···O hydrogen bond of length 2.24(8) Å, and neighboring pyramidal trinuclear [(µ3-Cl){HgSC(CH3)2CH(NH3)- COO_{3} ²⁺ moieties are further connected by symmetrical chloride bridges [Hg-Cl = 3.06(2) Å; $Hg-Cl-Hg'' = 79.6(7)^{\circ}$ to form a three-dimensional network. The voids in the lattice are filled by hydronium ions and novel planar cyclic hydrogen-bonded $(H_2 O \cdot C\Gamma)_3$ rings of edge $O - H \cdot \cdot \cdot Cl = 2.46(4)$ Å.

Introduction

The chemical behaviour of the element mercury differs significantly from those of its congeners zinc and cadmium, particularly as it relates to biochemistry and toxicology. In contrast to zinc, which is an important biological metal, mercury acts as a protein inhibitor in biological systems. Mercuric chloride, one of the important inorganic mercury pollutants, is a largely covalent molecule which remains undissociated in aqueous solution. Its interaction with sulfhydryl amino acids is in most cases the key to the biochemical action of inorganic mercury. While L-cysteine is usually used as a

model for the behaviour of cysteinyl-containing proteins towards mercurials, D-penicillamine and its derivatives constitute one of the most effective classes of antidotes for mercury poisoning. X-Ray structural studies have demonstrated linear or tetrahedral coordination of sulfhydryl groups or chloride ions around Hg²⁺ ions in the 1:2 and 1:1 HgCl₂/ L-cysteine complexes [1], linear S-Hg-S coordination in the 1:2 HgCl₂/D,L-penicillamine complex, and diagonal Cl-Hg-S coordination in the 2:1 HgCl₂/ D,L-penicillamine complex [2]. In the last complex, $(HgCl_2)_2$ [SC(CH₃)₂CH(NH₃)COOH] · 2H₂O, the [Cl-Hg-SC(CH₃)₂CH(NH₃)COOH] ⁺ and [HgCl₃]⁻ the structural units are interlinked by a weak thioethertype bridge from the penicillamine ligand. In solution, the complex readily dissociates into [Cl-Hg-S]⁺ and [HgCl₃]⁻ ions [3]. Since the existence of a [ClHgSC(CH₃)₂CH(NH₃)COOH]⁺ cation, and of similar species with different sulfhydryl amino acids like homocysteine and cysteine, has been postulated in one possible mechanism for environmental methylmercury formation [4, 5], it is desirable to isolate this 1:1 HgCl₂/penicillamine complex for structural analysis. In the present work, we report the synthesis and structural characterisation of a complex of stoichiometry $C_{30}H_{72}Cl_8Hg_6N_6O_{17}S_6$, of which a preliminary study has appeared as part of a doctoral dissertation [6].

Experimental

Preparation

Mercuric chloride (7 mmol in 20 ml ethanol) was added to D-penicillamine free base (7 mmol in 20 ml water) with stirring for 10 minutes. The resulting solution was then allowed to evaporate slowly at room temperature. Colourless rhomboidal dodecahedral crystals were obtained. Similar prepara-

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TABLE I. Data Collection and Processing Parameters.

Molecular formula	$C_{30}H_{72}Cl_8Hg_6N_6O_{17}S_6$		
Formula weight	2468.49		
Cell constants	a = 18.679(5) A, $V = 6518(6)$ A ³		
Density	$D_c = 2.516 \ (Z = 4), D_m = 2.60 \ \mathrm{g \ cm^{-3}}$		
Space group	P4 ₁ 32		
Crystal size	average radius $R = 0.04$ cm		
Radiation	graphite monochromatized		
	Mo- K_{α} ($\lambda = 0.71069$ A)		
μ (Mo-K $_{\alpha}$)	146.67 cm^{-1}		
Mean μR	2.7		
Transmission factors	0.004-0.013		
Scan type and speed	$\theta - 2\theta$; 2.55 to 14.55 deg min ⁻¹		
Scan range (2θ)	0.7° below $K_{\alpha 1}$ to 0.7° above $K_{\alpha 2}$		
Collection range	h, k, l ($h \le k \le l$); $2\theta < 45^{\circ}$		
Background counting	stationary counts for 0.4 of scan time at each end of scan		
Temperature	22 ± 1 °C		
Standard reflections	333, 333, 333		
Number of reflections collected between checks	125		
Unique data measured	696		
Observed data, n	443 (>3 <i>a</i> _F)		
Number of variables, p	72		
$R_F = \Sigma \ F_o - F_c / \Sigma F_o $	0.086		
$R_{w} = \left[\sum w (F_{o} - F_{c})^{2} / \sum w F_{o} ^{2} \right]^{\frac{1}{2}}$	0.101		
Weighting scheme	$w = \left[\sigma_{F}^{2}(F_{O}) + 0.003 F_{O} ^{2}\right]^{-1}$		
Largest shift in the last cycle	0.01		
Goodness of fit, S	1.305		
F(000)	4766.67 eA^{-3}		

tion using D,L-penicillamine yielded crystals giving the same X-ray diffraction pattern, indicating that identical or enantiomorphous crystals were obtained. *Anal.* Calcd. for $C_{30}H_{72}Cl_8Hg_6N_6O_{17}S_6$: C, 14.08; H, 3.23; N, 3.28. Found: C, 14.60; H, 3.08; N, 3.43%.

The bromo-analog was prepared in the same way by the reaction of mercuric bromide with D,L-penicillamine or D-penicillamine. X-Ray photographs confirmed that it is iso-structural with the chloroanalog. *Anal.* Calcd. for $C_{30}H_{72}Br_8Hg_6N_6O_{17}S_6$: C, 12.36; H, 2.84; N, 2.88; Br, 21.94. Found: C, 11.87; H, 2.74; N, 2.92; Br, 22.04%.

Crystallographic Study

Preliminary Weissenberg and precession photographs suggested cubic space group $P4_132$ (O^7), No. 213; systematic absences: h00, $h \neq 4n$) which was further confirmed by the Harker peaks on a Patterson map [6]. Another fresh crystal was mounted on a Nicolet R3m automated four-circle diffractometer. Accurate unit-cell dimensions were determined from a least-squares fit of 22 reflections $(11^{\circ} < 2\theta < 19^{\circ}; \text{ Mo-K}\alpha \text{ radiation})$. The reflection intensities declined rapidly with increasing Bragg angle, and data were collected in the range $0^{\circ} < 2\theta < 45^{\circ}$. Three standards monitored every 125 reflections showed only small random fluctuations within $\pm 1\%$, and linear scaling was applied. In addition to Lorentz and polarization factors, absorption correction was also applied using an empirical method based on a pseudo-ellipsoidal treatment of intensity measurements of selected strong reflections at different azimuthal angles. Other data collection and processing details are listed in Table I.

A sharpened Patterson map readily revealed the position of the Hg atom in the molecule. Other nonhydrogen atoms were located on subsequent difference Fourier maps. Anisotropic thermal parameters for the Hg, Cl, and S atoms and isotropic thermal parameters for the O, N, and C atoms were varied. Blocked-cascade least-squares refinement converged at $R_F = 0.086$ for 443 observed reflections. In view of the serious X-ray absorption of the crystal, no attempt was made to assign positions for

Atom	Site Symmetry	<i>x</i>	У	Z	$U_{\rm iso}/U_{\rm ec}$
Hg	1	78.0(2)	109.2(2)	226.6(2)	8.1(3)
Cl(1)	2	125	-46(1)	204(1)	10(3)
Cl(2)	2	125	471(1)	721(1)	7(2)
C1(3)	3	173(1)	173	173	7(1)
5	1	-4(1)	54(1)	303(1)	6(1)
D(1)	1	-189(3)	225(3)	375(3)	10(2)
D(2)	1	-93(2)	281(2)	398(2)	6(1)
D(3)	2	125	284(3)	534(3)	12(3)
D(4)	3	53(2)	53	53	4(2)
N	1	-140(3)	138(3)	269(3)	8(2)
C(1)	1	-122(4)	228(3)	369(3)	9(2)
C(2)	1	-85(3)	178(3)	315(3)	3(1)
C(3)	1	-42(3)	121(3)	357(3)	5(2)
C(4)	1	23(3)	163(4)	404(3)	6(2)
C(5)	1	-90(4)	80(5)	414(4)	10(3)

TABLE II. Fractional Coordinates $(\times 10^{-3})$ and Isotropic Thermal Parameters^a $(A^2 \times 10^{-2})$ of all Non-hydrogen Atoms.

^aForm of isotropic temperature factor: $\exp(-8\pi^2 U_{iso}\sin^2\theta/\lambda^2)$. U_{eq} calculated as one-third of the orthogonalized U matrix.

TABLE III. Bond Distances (A) and Angles (deg).

[Cl{HgSC(CH ₃) ₂ ($CH(NH_3)COO_3]^2$	trinuclear moiety			
Hg-Cl(3)	2.37(1)	Hg–Cl(3)–Hg'	98.5(9)	C(1)-C(2)-C(3)	109(2)
Hg-S	2.32(2)	Cl(3)-Hg-S	167.2(9)	O(1)-C(1)-O(2)	116(6)
S-C(3)	1.77(6)	Hg-S-C(3)	108(2)	O(1) - C(1) - C(2)	119(6)
C(3)-C(2)	1.54(8)	S-C(3)-C(2)	114(4)	O(2) - C(1) - C(2)	124(6)
C(3)-C(4)	1.68(9)	S-C(3)-C(4)	110(4)	$Hg \cdots Cl(1) \cdots Hg''$	79.6(7)
C(3)-C(5)	1.59(10)	S-C(3)-C(5)	105(4)	$Cl(1)\cdots Hg-Cl(3)$	101.9(3)
C(2)-N	1.54(8)	C(2)-C(3)-C(4)	109(5)	$Cl(1)\cdots Hg-S$	81.6(5)
C(2) - C(1)	1.54(8)	C(2) - C(3) - C(5)	112(5)		
C(1)-O(1)	1.25(9)	C(4)-C(3)-C(5)	106(5)		
C(1)-O(2)	1.25(8)	N-C(2)-C(1)	111(5)		
$Hg \cdots Cl(1)$	3.06(2)	N-C(2)-C(3)	108(4)		
Weak interactions	and hydrogen bor	nding			
Hg···O(4)	3.44(5)				
O(2)···O(2')	2.24(8)	$C(1) - O(2) \cdots O(2')$	119(4)		
$Cl(2)\cdots O(3)$	2.46(4)	$O(3) \cdots Cl(2) \cdots O(3')$	115(4)	$Cl(2)\cdots O(3)\cdots Cl(2')$	125(4)

the hydrogen atoms. The final difference map contained residual peaks in the neighbourhood of the Hg and S atoms, with extrema in the range 1.0 to $-1.1 \text{ e}^{A^{-3}}$.

All computations were performed on a Data General Corporation Nova 3/12 minicomputer with the SHELXTL program package [7, 8]. Analytical expressions [9] of neutral atomic scattering factors [10] were employed, and anomalous dispersion corrections [11] were incorporated. The final positional and isotropic thermal parameters are given in Table II. Tables of anisotropic thermal parameters for the Hg, Cl, and S atoms and observed and calculated structure factors have been deposited at the Editorial Office in Padua.

Results and Discussion

For structural reasons the title compound is best formulated as $2[(\mu_3-\text{Cl}){\text{Hg}(\text{penH})}_3]^{2+}\cdot 2H_3O^+$ $3(\mu_2-\text{Cl})^- \cdot (H_2O^+\text{Cl}^-)_3$, where penH designates the penicillaminato ligand [SC(CH₃)₂CH(NH₃)COO]⁻. Bond distances and angles are given in Table III in accordance with the atomic labelling in Fig. 1.

In the crystal lattice three crystallographically equivalent $[Hg(penH)]^+$ units are symmetrically linked by a triply bridging chloride ion [strong Hg-Cl(3) bond of lengths 2.37(1) Å; Hg-Cl(3)-Hg' = 98.5(9)°]. Each of these pyramidal trinuclear $[(\mu_3-Cl){Hg(penH)}_3]^{2+}$ moieties (Fig. 1) interacts with three others through moderately strong sym-

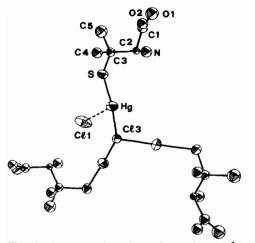


Fig. 1. A perspective view of the $[(\mu_3-Cl){HgSC(CH_3)_2-CH(NH_3)COO}_3]^{2+}$ unit, including one of the doubly bridging chloride ligands. The atom labelling scheme corresponds to that of Table II.

metrical chloride bridges [Hg-Cl(1) = 3.06(2) Å; $Hg-Cl(1)-Hg'' = 79.6(7)^{\circ}$, forming a threedimensional network. The carboxylate groups of two neighboring penicillamine ligands are strongly linked via a symmetrical O····H····O hydrogen bond, whose length of 2.24(8) Å is comparable to those reported in the literature [1, 12–14]. The hydrogen bonding scheme is also consistent with the observed carboxyl vibration modes $[\nu(C=0) \ 1740 \ (s, br); \ \nu(C=0)$ 1300 cm^{-1} (m, br)]. The protons involved in this type of hydrogen bonding come from half of the amino groups. Thus on the average, 21/2 protons reside on each amino group in the unit cell. This kind of disordering is supported by the observed vibrational frequencies $[\delta_{as}(-NH_3^+) 1570 \text{ (w, br)};$ $\delta_{s}(-NH_{3}^{+})$ 1480 cm⁻¹ (w, sh)] [6] which are lower than those for penicillamine hydrochloride $[\delta_{as}]$ $(-NH_3^+)$ 1624(m); $\delta_s(-NH_3^+)$ 1520 cm⁻¹ (s)] [15] but higher than those for the N-bonded complexes $(CH_3Hg)_2(pen) [\delta_{as}(-NH_2^+-) 1586(vs); \delta_s(-NH_2^+-)]$ 1230 cm⁻¹ (w, sh)] [16] and $CH_3Hg(D,L$ -methionine) $[\delta_{as}(-NH_2^{+})] = 1590 \text{ (vs, br)}; \ \delta_{s}(-NH_2^{+})$ $1250 \text{ cm}^{-1} \text{ (m, br)} [17].$

Hydrogen bonding between Cl(2) and the water molecules of crystallisation forms planar six-membered $(H_2O \cdot Cl^-)_3$ rings of edge $O(3) - H \cdot \cdot \cdot Cl(2) =$ 2.46(4) Å. These strongly hydrogen bonded rings fill the voids of the three-dimensional network constructed by [Hg(penH)]⁺, Cl(1), and Cl(3) (molecular packing shown in Fig. 2), with no interatomic contact less than 4 Å between the two components. To our knowledge the $(H_2O \cdot C\Gamma)_3$ ring is novel and constitutes a higher homolog of the recently discovered planar cyclic hydrogen-bonded (H₂O·Br⁻)₂ system in crystalline $[(CH_2)_6N_4CH_3]Br \cdot H_2O$ [18]. Atom O(4) occupies a site of symmetry 3 and only weakly interacts with Hg [Hg···O(4) = 3.44(5) Å]. To preserve overall electroneutrality in the crystal, O(4) is identified with the hydronium (or oxonium) ion H_3O^+ .

The coordination around the Hg atom is mainly diagonal with significant distortion from linearity $[Cl(3)-Hg-S = 167.2(9)^{\circ}]$. The strong Hg-X bonds [Hg-Cl(3) = 2.37(1) and Hg-S = 2.32(2) Å] are comparable to those in similar diagonal S-Hg-X (X = S, Cl) systems such as the L-cysteine complex $Hg(cystH_2)(cystH)Cl \cdot \frac{1}{2}H_2O$ [Hg-S = 2.353(3) and 2.329(5) Å] [1], where cystH stands for the Lcysteinato ligand [SCH₂CH(NH₃)COO]⁻⁻, or the D,L-penicillamine complexes $[Hg(penH_2)_2Cl]Cl \cdot H_2O$ $[Hg-S = 2.335(5) \text{ and } 2.357(4) \text{ Å}] \text{ and } (HgCl_2)_2$ - $(\text{penH}_2) \cdot 2\text{H}_2\text{O}$ [Hg(2)-S = 2.356(5) and Hg(2)-Cl(4) = 2.356(5) Å] [2]. All these Hg-X bonds are distinctly shorter than corresponding bonds in the tetrahedrally coordinated L-cysteine complex $HgCl_2(cystH_2)$ [Hg-S = 2.453(4) and 2.490(4); Hg-Cl = 2.582(4) and 2.645(5) Å] [1].

The linearity of the S-Hg-X units is susceptible to distortion by weaker interactions with other ions. The Cl(3)-Hg-S angle in the present 1:1 complex $2[Cl{Hg(penH)}_3] \cdot 3Cl \cdot 2H_3O \cdot (H_2O \cdot Cl)_3$ deviates by 12.8° from 180° because of the participation of Cl(1) in the primary coordination shell around Hg. While the chloride ions are excluded from the primary coordination shell in the 1:2 HgCl₂/cystH₂ complex [Hg···Cl = 3.232(5) Å], they interact strongly with Hg atoms in the 1:1 complex [1].

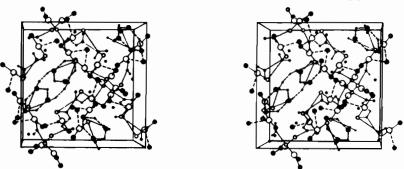


Fig. 2. Packing and hydrogen bonding in $2[(\mu_3-Cl){Hg(penH)}_3] \cdot 3(\mu_2-Cl) \cdot 2(H_3O) \cdot (H_2O-Cl)_3$. For clarity, the $-(CH_3)_2CH \cdot (NH_3)$ - fragment is represented by a line linking S with C(1) of the carboxylate group.

TABLE IV. Torsion Angles (deg) in the D-penicillamine Skeleton.^a

$O(1) = C(1) - C(2) - N, \psi_{T}^{1}$	9(8)	C(1)-C(2)-C(3)-S	174(4)
O(1)=C(1)-C(2)-C(3)	-110(7)	C(1)-C(2)-C(3)-C(4)	-63(6)
$O(2)-C(1)-C(2)-N, \psi_{T}^{2}$	-158(6)	C(1)-C(2)-C(3)-C(5)	54(7)
O(2)-C(1)-C(2)-C(3)	83(7)	C(2)-C(3)-S-Hg	63(4)
$N-C(2)-C(3)-S, \chi^{1}$	54(6)	C(4) - C(3) - S - Hg	-59(4)
N-C(2)-C(3)-C(4)	176(4)	C(5)-C(3)-S-Hg	-173(4)
N-C(2)-C(3)-C(5)	-66(6)	· _	

^aNotations ψ_{T}^{1} , ψ_{T}^{2} , and χ^{1} follow the convention in: IUPAC-IUB Commission on Biochemical Nomenclature, *J. Mol. Biol.*, 52, 1 (1970).

The same trend is also found for the $HgCl_2/penH_2$ complex series but with even stronger metal-chloride interactions. Hence the participation of chloride ions in the primary coordination shell increases as the $Hg/penH_2$ ratio in the compound decreases. The 1:2 complex [Hg(penH₂)₂Cl]Cl·H₂O consists mainly of two-coordinate S-Hg-S with a moderately strong Hg-Cl bond [Hg-Cl(1) = 2.850(5); also weakinteraction $\operatorname{Hg} \cdots \operatorname{Cl}(1') = 3.323(5)$ Å] [2]. One of these two Hg-S bonds is replaced by a very strong Hg-Cl bond [2.37(1) Å; Cl triply bridging] with retention of the moderately strong Hg-Cl bond [3.06(2) Å] in the present 1:1 complex, whereas the $[ClHgSC(CH_3)_2CH(NH_3)COOH]^+$ moiety in the 2:1 complex (HgCl₂)₂(penH₂)·2H₂O consists of a distorted linear S-Hg-Cl system with four bridging chloride interactions [Hg-Cl in the range 2.780(5)-3.429(5) Å] [2]. Thus from the structural point of view the conversion of one HgCl₂/penH₂ complex to another can be rationalised in the following way. Replacement of a penicillamine ligand in the 1:2 complex with a chloride ion gives the 1:1 complex, and the 2:1 complex can be regarded as an adduct of HgCl₂ with the 1:1 complex. In fact, an nmr study has provided evidence for the interconversion of $Hg^{2+}/penH_2$ complexes in solution [6].

Due to the weak diffracting power of crystalline $2[Cl{Hg(penH)}_{3}] \cdot 3Cl \cdot 2H_{3}O \cdot (H_{2}O \cdot Cl)_{3}$ and the predominate scattering of the heavy Hg atom, it has not been possible to determine the bond distances for the penicillamine skeleton with good precision. They are all in reasonable agreement, within experimental errors, with corresponding values found in the free base [15], as well as in its hydrochloride [19] or complexes like $[Hg(penH_2)_2Cl]Cl \cdot H_2O$, $(HgCl_2)_2(penH_2) \cdot 2H_2O$ [2, 3], and $[Co(L-histidine) \cdot 2H_2O$ (D-penH)]·H₂O [20]. However, the penicillamine ligand adopts different conformations in these compounds. Torsion angles of interest for the Dpenicillamine skeleton in the present complex is listed in Table IV. The torsion angle C(1)-C(2)-C(3)-S = $174(4)^{\circ}$ indicates that the bulky sulfur atom is antiperiplanar to the carboxyl terminus. This is a characteristic conformation for compounds with protonated carboxylate groups. In those compounds like meso-penicillamine [15], CH₃Hg(penH)·H₂O, and $(CH_3Hg)_2(pen)$ [16], where intramolecular interactions bend the skeleton to form six-membered rings [O-C(1)-C(2)-C(3)-S-M, M = Hg or H], the bulky sulfur atoms are ±synclinal to the deprotonated carboxylate groups and are located between the comparatively large amino and carboxyl groups, resulting in higher conformation energies. Further twisting of the penicillamine skeleton is observed for tridentate ligands such as in the S,N,O-bonded complex [Co(L-histidine)(D-penH)]·H₂O [20]. Another intramolecular interaction in the present complex, namely the possible hydrogen bonding between the amino and carboxyl groups $[N \cdots O(1) 2.71(8) \text{ Å}]$, leads to torsion angles $O(1)=C(1)-C(2)-N(\psi_T^1)$ and O(2)- $C(1)-C(2)-N(\psi_{T}^{2})$ of 9(8) and -158(6)° respectively, in contrast to those [-21.2 to -23.6 and 156.4 to 164.5° respectively] for L-penH₂·HCl· H_2O [19], [Hg(penH₂)₂Cl]Cl·H₂O, and (HgCl₂)₂- $(penH_2) \cdot 2H_2O$ [2, 3], where the amino groups are in the proximity of the carbonyl oxygen atoms.

It has been found that the complex $[Ni(D-pen)_2]^{2-}$ is thermodynamically more stable than $[Ni(D-pen)(L-pen)]^{2-}$ [21]. The enantiomorphous crystallisation observed in the present 1:1 HgCl₂/penH₂ system provides an interesting example of stereochemical selectivity, which may be contrasted wih the reaction of CoCl₂·2H₂O with D,L-histidine followed by addition of D,L-penicillamine to yield crystalline [Co(L-his)(D-pen)] [Co(D-his)(L-pen)] · 2H₂O [22].

Acknowledgement

The authors wish to thank Professors A. J. Carty and C. Chieh (University of Waterloo, Ontario, Canada) for their interest and advice.

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