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 ${(CH_3)_2SO}_2CuBr_2$ is isomorphous with the corresponding chloride complex. It belongs to the orthorhombic space group Pnma and contains 4 formula groups per unit cell. The dimensions of the unit cell at ca. 295K are a = 8.257(2)Å, b = 11.772-(2)Å, c = 11.749(2)Å. The copper ion assumes an intermediate five-coordinate geometry with three bromine atoms equatorial and two oxygen atoms axial. The equatorial angles are $Br(1)-Cu-Br(2) = 146.13^{\circ}$ and $Br(2)-Cu-Br(2) = 110.21^{\circ}$, and the axial angle O(1)-Cu-O(1') is 173.99°. The bond lengths are Cu-Br(1) = 2.451(4)Å, Cu-Br(2) = 2.429(4)Å, Cu-Br(2') = 2.777(4)Å and Cu-O = 1.962(9)Å. The copper coordination geometry is largely determined by packing of the DMSO groups.

Introduction

As part of a program of structural characterization of copper(II) bromide salts, we have undertaken the crystal structure determination of $\text{CuBr}_2 \cdot 2\text{DMSO}$. While the structural literature of copper(II) chloride salts is profuse [1], a relatively small number of bromide complexes have been the subject of structural characterization. The distorted (compressed) tetrahedral geometry, as defined in Cs₂CuBr₄ [2], has an average Cu-Br distance of 2.35Å with bond angles near 100° and 125°. This distance is longer (2.39Å) in the more distorted anion present in (enH₂Br)₂CuBr₄ [3], indicating this distance is sensitive to distortion. In the distorted (elongated) octahedral geometry, as exemplified in CsCuBr₃ [2], the average axial Cu-Br distance is 2.45Å, although the individual distances range from 2.34Å to 2.52Å. The axial distances are 2.87 and 2.96Å. For the trigonal bipyramidal geometry, the axial Cu–Br bond length is 2.45 Å while the equatorial distance is 2.52Å [4, 5]. The bis(dimethylsulfoxide) complex of CuBr₂ was chosen for study, since the isomorphous chloride had a coordination geometry intermediate been square pyramidal and trigonal bipyramidal [6]. This, hopefully, would allow the estimation of steric effects in these copper(II) salts which fall in this gray area of five-coordinate geometry intermediate between C_{4v} and D_{3d} symmetry.

Preparation and Data Collection

Small dark brown platelets of $CuBr_2(DMSO)_2$ can be prepared by reacting ethanolic solutions of $CuBr_2$ and DMSO and allowing the slow evaporation of the solvent [7].

Examination of Weissenberg, X-ray diffraction, photographs of a wedge shaped crystal (ca. 0.18 × 0.16 × 0.15 mm × 0.01 mm thick) showed it to belong to the orthorhombic class having mmm diffraction symmetry. The lattice constants measured with Zr filtered MoK_{α} radiation using a Picker diffractometer and single crystal orienter were found to be a =8.257(2)Å, b = 11.772(2)Å, and c = 11.749(2)Å. Systematic absences for 0kl when k + 1 = 2n + 1 and for hk0 when h = 2n + 1 suggested that the space group was the same as that found for the corresponding chloride, namely Pnma.

TABLE I. Final Positional and Thermal Parameters^a for CuBr₂·2DMSO.

	x	у	z	β_{11}	β22	β ₃₃	β ₁₂	β ₁₃	β ₂₃
Cu	3157(3)	2500	3672(2)	107(5)	18(2)	48(2)	0	110(3)	0
Br(1)	4933(3)	2500	2000(2)	104(4)	43(2)	67(2)	0	340(3)	0
Br(2)	3011(4)	2500	5737(2)	192(6)	42(2)	48(2)	0	-8(3)	0
S	1834(5)	4805(3)	4264(3)	107(7)	25(3)	51(3)	0(4)	19(5)	4(3)
õ	3123(13)	4165(7)	3587(8)	116(19)	25(7)	55(9)	20(11)	43(12)	-17(7)
cm	3004(26)	5798(15)	5074(20)	240(47)	41(15)	173(26)	-2(23)	-64(32)	-44(17)
C(2)	932(26)	5734(16)	3231(15)	255(48)	57(16)	90(19)	57(23)	-29(26)	2(14)

^a Parameters multiplied by 10⁴.

	Distance			Angle	
	Cl	Br		Cl	Br
Cooper Coordinat	ion Geometry				
Cu-X(1)	2.290(2)	2.451(4)	X(1)-Cu- $X(2)$	146.1(1)	146.1(2)
Cu-X(2)	2.284(2)	2.429(4)	X(1)-Cu-O	88.7(1)	88.2(3)
Cu-O	1.955(4)	1.962(9)	X(2) = Cu = O	92.9(2)	92.9(3)
$C_{u} = \mathbf{Y}(1)^{\mathbf{a}}$	2 702(2)	2 777(4)	X(1) = Cu = O $X(1) = Cu = X(1)^{a}$	1127(5)	110 2(1)
Cu-A(1)	2.702(2)	2.777(4)	$X(2) - Cu - X(1)^{a}$	101.2(1)	103.7(1)
			0-Cu-O	173.0(3)	173.9(6)
			Cu-X(1) ^a -Cu ^a	144.6(1)	
DMSO Geometry					
SO	1.531(4)	1.53(1)	0-S-C(1)	104.7(3)	102.9(8)
S-C(1)	1.771(7)	1.79(2)	O-S-C(2)	103.9(3)	103.7(7)
S-C(2)	1.765(7)	1.79(2)	C(1)-S-C(2)	100.4(3)	100.7(9)
			Cu-O-S	118.2(2)	118.5(6)
Intermolecular No	on-bonded Interactions	8			
C(1)–Br (1) ^b		3.87(2)			
C(1)-Br(2)		3.96(2)			
C(1)–Br(2) ^c		3.97(2)			
$C(2)-Br(2)^d$		3.70(2)			
C(2)-Br(1) ^a		3.90(2)			
C(1)-C(1)		3.80(4)			
Coordination Geo	metry Non-bonded In	teractions			
X(1)-X(2)	4.376(1)	4.668(3)			
$X(1) - X(1)^{a}$	4.161(1)	4.292(3)			
$X(2)-X(1)^{a}$	3.862(1)	4.070(3)			
X(1)-0		3.09(1)			
X(2)-O		3.20(1)			
X(1) ^a –O		3.35(1)			

TABLE II. Interatomic Distance and Angles. Standard Deviation of Last Digit in Parentheses.

^aAtom transformed by $-\frac{1}{2} + x$, $\frac{1}{2} - y$, $\frac{1}{2} - z$. ^bAtom transformed by $\frac{1}{2} - x$, 1 - y, $\frac{1}{2} + z$. ^cAtom transformed by 1 - x, $\frac{1}{2} + y$, 1 - z. ^dAtom transformed by $\frac{1}{2} - x$, 1 - y, $-\frac{1}{2} + z$.

A total of 1024 independent reflections were measured out to $2\theta = 50^{\circ}$ with Zr filtered MoK_a radiation. A θ -2 θ step scan of 2.0° degrees was used with 20 steps/degree and 4 sec/step. Background measurements were made before and after each scan. Absorption corrections ($\mu = 43.9 \text{ cm}^{-1}$) were made. The standard deviation of each reflection was calculated by the expression σ^2 (I) = TC + BC + (0.03)²I_o² where TC = total counts, BC = background counts, and I_o = TC - BC.

The structure was determined by isomorphous substitution of Br for Cl in the structure determined for CuCl₂(DMSO)₂. Refinement converged quickly to a final weighted R factor $[\omega_{bk1} = 1/\sigma^2(F)]$ of 0.088 and a conventional R factor of 0.083 for the reflections with $F > 3\sigma(F)$. No attempt was made to find the hydrogen atoms. The final difference map showed no unusual features. The final parameters are reported in Table I and Table II gives pertinent interatomic distances and angles for the Br salt and the isomorphous Cl salt. An illustration of the molecule is given in Fig. 1 while the packing is illustrated in Fig. 2. Computer programs used were part of a local program library [8, 9] containing modified versions of ORFLS3 [10], ALFF [11], ORFFE3 [12], and ORTEP [13].

Discussion

As expected from the isomorphism of $CuCl_2$. 2DMSO and $CuBr_2$.DMSO, the structures of the two salts are virtually identical. In fact, with the exception of the Cu-X (X = Cl, Br) *distances*, the isomorphism is so complete it is unnecessary to comment on many features of the structure.

In the analysis of the properties of $CuCl_2 \cdot 2DMSO$ [6], it was shown that it was more appropriate to

TABLE III. Observed and Calculated Structure Factors for $CuBr_2 \cdot 2DMSO$. Columns are k, l, $10|F_0|$ and $10F_c$.

(Continued overleaf)

TABLE III. (Continued)

223333333333333333333	12 13 4 5 6 7 8 9 10	49 -237 254 -308 1500-1350 0 38 893 847 474 457 929 -966 123 116 320 361 0 19 259 239 95 86	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	12 3 12 4 4 12 5 4 12 6 2 13 0 2 13 1 2 13 3 3 13 4 2 H = 0 1 12	0 -11 10 406 421 10 402 -425 10 205 163 11 264 251 11 250 -117 11 323 262 11 3 11 3 11 279-1274 12	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	9 4 9 5 9 6 9 7 9 8 10 0 10 1 10 2 10 3 10 4 10 5 10 6	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$ \begin{array}{cccccccccccccccccccccccccccccccccccc$	$\begin{array}{cccccccccccccccccccccccccccccccccccc$	$\begin{array}{cccccccccccccccccccccccccccccccccccc$
3	11	42 68	2 3 294 -295	0 1 14	219-1219 12	2 205 -21	10 0	0 18 0	3 441 -408 3	3 157 -37	
э	12	63 -139	2 4 887 -886	0 2 1	148 -90 12	3 524 499	10 7	0 -131 0	4 337 -352 3	4 545 519	



Fig. 1. Coordination sphere of copper(II) in CuBr₂·2DMSO.

consider the coordination geometry as distorted square pyramidal than distorted trigonal bipyramidal. Examination of bond distances in the bromide analog would indicate it is again more appropriate to describe the geometry as distorted square pyramidal. The two shorter Cu-Br distances (2.43, 2.45Å) are comparable to the equatorial distances for the distorted octahedral geometry in CsCuBr₃ (2.45Å average) [2]. The equatorial Cu-Br distances for the trigonal bipyramidal geometry (2.52Å) are distinctly longer [4, 5]. Thus, we envisage the structure as derived from the severe distortion of a planar fourcoordinate species (two bromine, two oxygen ligands) by the presence of a fifth axial ligand.

It is instructive to make a detailed comparison of the interatomic distances in the copper chromophores of the two DMSO complexes. The axial Cu-Br distance is 2.777(4)Å, approximately 0.34Å longer than the other two Cu-Br distances. The opposite bond angle is $146.13(2)^{\circ}$. In the chloride salt, the axial Cu-Cl bond is 0.42Å longer than equatorial bonds while the bond angle is identical. Thus the axial bond only elongated 0.08Å. The change in the short bond Cu-X bond lengths between the chloride and bromide salt is roughly 0.15Å, corresponding precisely to the change in ionic radii of the two halide ions. Thus the only steric compression in the chromophore upon change from bromide to chloride is in the axial Cu-X distance. This is born out by examination of the non-bonded interactions in the coordination sphere. The only short contacts are the X-O inter-



Fig. 2. Stereographic view of the crystal structure of CuBr₂·2DMSO as viewed from near the b-axis.

actions since the X-X interactions are all greater than the sum of the van der Waals radii. The X_1 -O and X_2 -O distances are less than the sum of the van der Waals radii since both ligands are strongly bonded to the copper. In the chloride salt, the Cl'₁-O distance is exactly equal to this sum, as pointed out previously. However, in the bromide salt, the Br'₁-O distance is 0.1Å shorter than the sum.

The source of this steric compression logically lies in crystal packing forces, implying that the coordination geometry is largely determined by *inter*molecular steric forces, rather than *intra*molecular forces. Indeed, examination of the packing of the chains clearly shows that the methyl groups and halides neatly interlock, fixing their locations. The copper geometry thus largely reflects these steric interactions, rather than the inherent stability of the observed geometry of the chromophore.

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