

In all the examples which follow, 'sharpened' $|F|^2$ values were used in the calculation of the Q -functions. $Q(X_0Z_0)$ was calculated using the relative coordinates of the eleven atoms in the plane of the purine residue in deoxyadenosine. The relative coordinates were those obtained from the determination of the molecular orientation (Watson, Sutor & Tollin, 1965). The space group is $P2_1$. The map obtained is shown in Fig. 1(a). The expected positions of peaks due to atoms separated by $b/2$ are marked in the map. The largest remaining peak is that which determines the position of the origin. The map of $Q(X_0Z_0)$ obtained using the relative coordinates for the atoms of the sugar residue in deoxyadenosine is shown in Fig. 1(b). In this case only eight out of a total of twenty heavy atoms in the molecules were used. The coordinates used were the final coordinates from the refined structure with an arbitrary change of origin to $x=0.2$, $z=0.3$. In this case there are no atoms separated by half in their fractional y coordinates.

The function $Q(Y_0)$ was calculated from relative coordinates for pyrimidine obtained by taking the final published coordinates (Wheatley, 1960) and giving them an arbitrary shift of origin to $x=-0.15$, $y=-0.3$. The space group is $Pna2_1$. $Q(Y_0)$ was used to define the position of the molecule relative to the a -glide plane. Only the fifty largest 'sharpened' $|F|^2$ values were used. The resulting map is shown in Fig. 2 with the origin shifted to $y=-\frac{1}{2}$ to allow for the fact that the a -glide occurs at $y=\frac{1}{2}$. The dotted vertical line represented the correct answer of $y=-0.3$. The results show that even with this small amount of data the origin position is well defined. It can also be seen that the determination of the y coordinate is independent of the fact that at this stage the x coordinate of the arbitrary origin is not known. In all these examples the error in determining the origin position was less than 0.05 \AA .

These Q -functions have also been used to determine the structure of 4-acetyl-2'-fluorobiphenyl (Tollin, Young & Sutherland, 1965).

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The crystal structure of bis(*N*-2-hydroxyethylsalicylaldiminato)copper (II). By E. R. BOYKO, *Chemistry Department, Providence College, Providence, Rhode Island, U.S.A.* and D. HALL, MARY E. KINLOCH and T. N. WATERS. *Chemistry Department, University of Auckland, New Zealand*

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There is considerable current interest in the crystal structures of *N*-substituted salicylaldiminato complexes of transition metals (see Table 5, Frasson, Panattoni & Sacconi, 1964; Fox, Lingafelter, Orioli & Sacconi, 1963; Wei, Stogsdill & Lingafelter, 1964; Cheeseman, Hall & Waters, 1965). Attention has been mainly focused on the dimensions and stereochemistry of the molecule, as in all but the simplest such molecules there appears to be little molecular interaction. The substituents in compounds studied to date have

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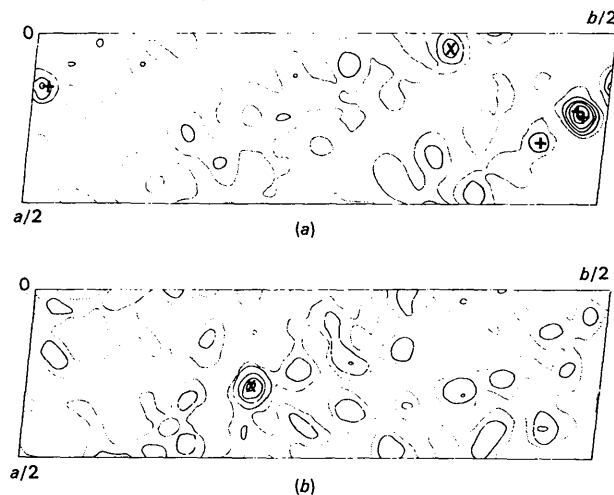


Fig. 1. $Q(X_0Z_0)$ for deoxyadenosine (a) using purine relative coordinates (b) using sugar relative coordinates. + indicates false peaks. x indicates expected positions.

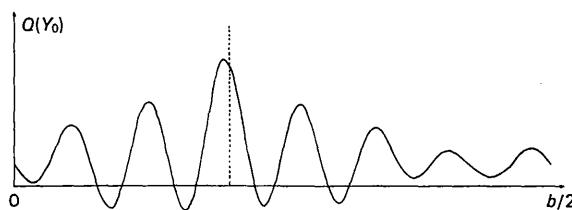


Fig. 2. $Q(Y_0)$ for pyrimidine. Dotted line indicates expected peak position.

been alkyl or phenyl groups, and it is of interest to consider the influence on such structures of groups with hydrogen-bonding potential. In this connection studies of the crystal structure of bis(*N*-2-hydroxyethylsalicylaldiminato)copper(II) were begun both at Providence and Auckland, the results from one of these being announced at the Rome Congress (Boyko, 1963). It transpired that the two investigations were similar in scope and achievement, and a joint publication has been decided upon.

Experimental

The compound was prepared by warming stoichiometric quantities of bisalicylaldehydocopper(II) and ethanolamine in 50% aqueous methanol, and recrystallized from ethanol as dark green needles, m.p. 176°. Chemical analysis

verified the resulting composition. The constants for the monoclinic unit cell were obtained from NaCl-calibrated Weissenberg photographs around *b* and *c* as $a = 18.66 \pm 0.04$, $b = 4.71 \pm 0.01$, $c = 19.99 \pm 0.04$ Å, $\beta = 97.8 \pm 0.1^\circ$. The density was measured by a gradient density procedure as 1.48 g.cm^{-3} , as compared with the calculated density of

Table 1. Atom coordinates and temperature factors

Atom in molecule A					Atom in molecule B				
Atom	<i>x</i>	<i>y</i>	<i>z</i>	<i>B</i>	Atom	<i>x</i>	<i>y</i>	<i>z</i>	<i>B</i>
Cu	0	0	0	4.5 Å ²	Cu	$\frac{1}{2}$	$\frac{1}{2}$	0	4.0 Å ²
O(1)	-0.0049	-0.1437	0.0863	3.8	O(1)	0.5732	0.5245	0.0761	3.4
N	-0.0885	0.2259	0.0066	4.3	N	0.4513	0.2052	0.0483	4.5
C(1)	-0.1301	0.2196	0.0532	3.8	C(1)	0.4766	0.1061	0.1086	3.0
C(2)	-0.1187	0.2596	0.1094	3.7	C(2)	0.4577	0.8014	-0.1517	3.9
C(3)	-0.0588	-0.1357	-0.1261	4.7	C(3)	0.4130	0.5886	-0.1358	3.3
C(4)	-0.0491	-0.3195	0.1841	5.2	C(4)	0.3478	0.5001	-0.1791	4.5
C(5)	-0.1058	-0.3223	-0.2250	4.9	C(5)	0.3299	0.6501	-0.2436	4.5
C(6)	-0.1675	-0.1508	0.2121	5.4	C(6)	0.3750	0.8516	-0.2626	4.5
C(7)	-0.1723	0.0281	0.1572	4.4	C(7)	0.4397	0.9318	-0.2173	4.1
C(8)	-0.1117	0.4511	-0.0472	4.3	C(8)	0.3854	0.0479	0.0743	3.7
C(9)	-0.1597	0.2882	-0.1104	3.1	C(9)	0.3154	0.2460	0.0110	3.7
O(2)	-0.2320	0.2293	-0.0930	3.8	O(2)	0.2970	0.2767	0.0801	4.2

Estimated standard errors

	<i>x</i>	<i>y</i>	<i>z</i>	<i>B</i> (Å ²)
Cu	—	—	—	0.2 Å ²
O	0.015 Å	0.020 Å	0.014 Å	0.4
N	0.018	0.024	0.017	0.4
C	0.021	0.031	0.020	0.55

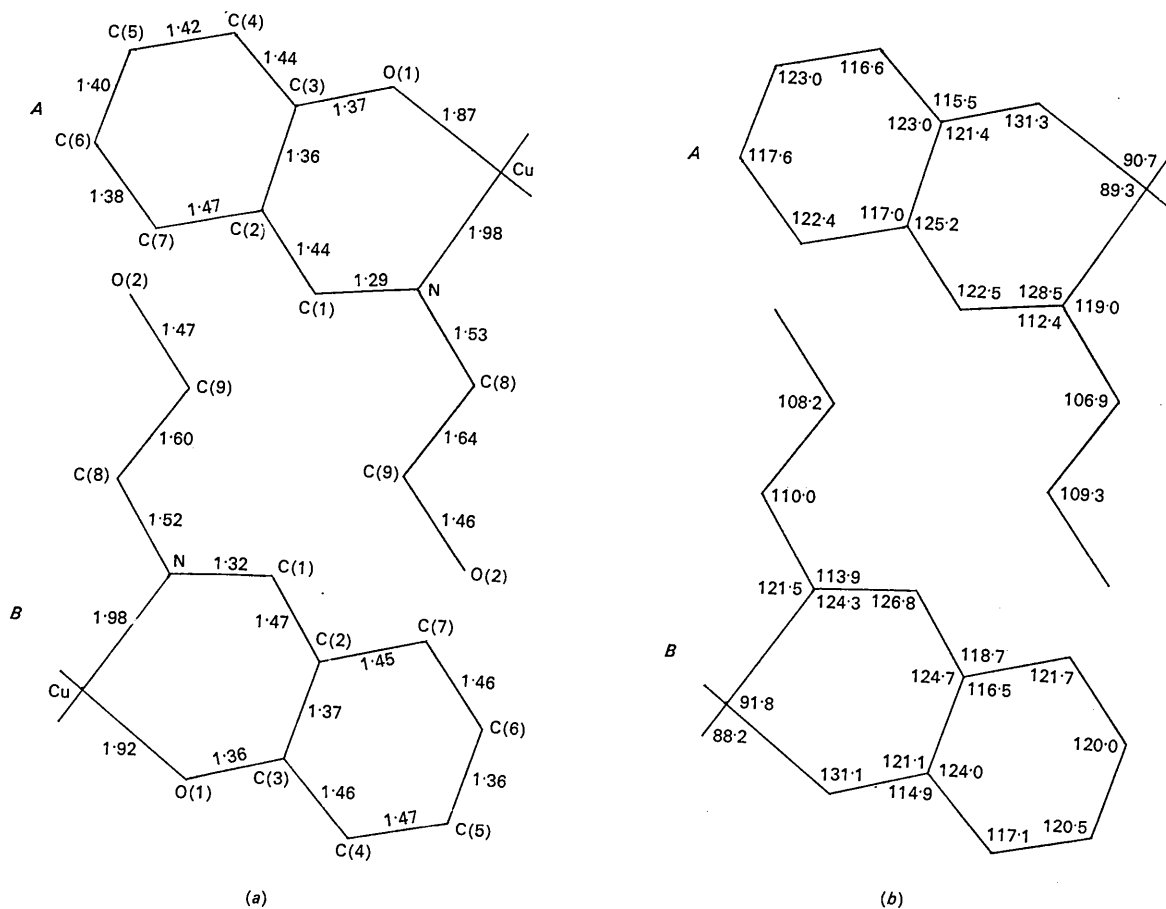


Fig. 1. (a) Bond lengths (Å). (b) Bond angles (°).

1.495 g.cm⁻³ for 4 molecules per unit cell. Systematic absences indicated the space group *P2₁/c*.

Intensities were measured visually from Weissenberg photographs of the *h0l*, *h1l* and *h2l* layers. The approximate dimensions of the crystal employed were 0.1 × 0.4 × 2 mm. No correction for absorption was made. The shape of the crystals precluded satisfactory photography other than about *b*, and the interlayer scaling constants were regarded as additional parameters during refinement. A comparison of the 600 observed structure factors from the two independent determinations showed no serious discrepancies and only one set (Providence) was employed in the refinement.

Structure determination

The Patterson function showed dominant maxima at positions ($\frac{1}{2}, \frac{1}{2}, 0$), ($\frac{3}{2}, 0, \frac{1}{2}$) and ($0, \frac{1}{2}, \frac{1}{2}$), whence it was deduced that the copper atoms either occupy general positions at ($\sim 0.25, \sim 0.00, \sim 0.25$) or the two independent sets of special positions at ($0, 0, 0$) and ($\frac{1}{2}, \frac{1}{2}, 0$). Heavy atom phased Fourier syntheses could only be interpreted on the latter assumption, *i.e.* that two independent centric molecules exist in the structure. Location of the light atoms and subsequent refinement followed conventionally, although the details of procedure in the two investigations differed somewhat. The final refinement was by a least-squares procedure in which atoms were permitted individual isotropic temperature factors. The atomic form factors employed were from Berghuis, Haanappel, Potters, Loopstra, MacGillavry & Veenendaal (1955) and the weighting factors used were $w = 1$ for $F_o \leq 30$, $w = 30/F_o$ for $F_o > 30$. Further refinement was attempted with anisotropic temperature factors, but little improvement resulted and the isotropic refinement was preferred for its smaller number of parameters. The overall reliability index (observed structure factors only) is 0.090; individual values for separate layers are 0.091, 0.087 and 0.095 for *h0l*, *h1l*, and *h2l* respectively. Atom coordinates and temperature factors are listed in Table 1, together with the estimated standard errors calculated from the least-squares residuals.

Discussion

The bond lengths and angles in the two independent centric molecules are shown in Fig. 1. In general they are similar, although the difference in the length of the bonds C(6)–C(7), 0.079 Å, and of Cu–O, 0.053 Å, would appear significant by normal criteria. It is unlikely that this is true, and more probable that the standard errors in Table 1 are underestimated. The excessively long values for both of the bonds C(8)–C(9) would support this conclusion. No further discussion of the individual dimensions is then profitable, other than to note the general similarity to those of the chemically related molecules bis-(*N*-methylsalicylaldiminato)copper(II) (Lingafelter, Simmons, Morosin, Scheringer & Freiburg, 1961) and bis-(*N*-phenylsalicylaldiminato)copper(II) (Wei *et al.*, 1964).

The two molecules in this study do, however, appear to differ meaningfully with respect to planarity. The mean planes (Schomaker, Waser, Marsh & Bergman, 1959) through the molecules (other than the side chains) and the mean planes through a benzene ring were calculated. For molecule *A* the approximation to overall planarity is poor; the benzene ring is effectively planar, however, and with the exception of the nitrogen (deviation 0.125 Å), the remaining atoms of the salicylaldimine group are coplanar

with it. The molecule is bent, or stepped, so that the planes of the two benzene rings are separated by 0.62 Å, this situation being similar to that observed in bis-(*N*-phenylsalicylaldiminato)copper(II). On the other hand molecule *B* can, with the sole exception of the oxygen atoms (deviation 0.092 Å), be described as coplanar within the accuracy

Table 2. Observed and calculated structure factors

	huo	h0l	h0k	hok	h0l	h0k	hok
2	1999 1828	0	1605 1565	0	822 772	0	127 220
3	216 -244	2	1252 1392	2	691 818	1	243 144
4	216 -217	4	1204 1222	3	736 450	2	1320 1335
5	927 8-2	4	862 1050	5	606 -563	3	169 -227
6	565 509	5	304 -303	6	780 806	4	213 340
7	537 506	6	603 648	7	786 -753	5	226 -260
8	508 459	7	515 420	8	900 852	6	679 366
9	508 557	8	1113 1097	9	172 70	7	231 -298
10	562 487	10	1126 1123	10	931 839	8	171 314
11	250 196	11	290 -376	11	194 156	9	184 -224
12	322 261	12	662 651	12	470 403	10	325 365
13	406 363	13	201 201	13	201 148	10	352 309
14	406 344	14	178 324	-1	869 884	14	218 159
15		15	208 178	-2	571 651	15	187 297
16		16	873 982	-3	675 667	-1	131 -195
17		17	1282 -1397	-5	183 -186	-2	496 -592
18		18	1138 1246	-6	440 544	-3	292 288
19		19	450 -399	-7	216 20	-4	532 77
20		20	194 295	-8	462 478	-6	786 767
21		21	262 280	-9	198 198	-7	169 190
22		22	706 654	-10	611 54	-8	290 350
23		23	381 373	-11	170 155	-9	599 -565
24		24	211 217	-12	344 350	-10	177 188
25		25	173 178	-14	504 390	-11	286 307
26		26	248 248	-16	242 229	-12	483 485
27		27	736 650	-17	574 -577	-12	278 283
28		28	418 325	-18	311 294	-18	269 228
29		29		-20	188 161	-20	280 281
30		30					
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100		100					

of these observations. Alternatively, consideration of the benzene rings alone would again suggest that it be described in terms of individually planar but not coplanar salicylaldimine groups, separated by 0.2 Å. The experimental accuracy is not sufficient to distinguish a planar molecule from one that is slightly stepped. These variations between *A* and *B* must presumably stem from packing effects, and would support the contention (Cheeseman, Hall & Waters, 1965) that the resistance of such molecules to deformation from overall planarity is rather less than has often been supposed.

The coordination is square planar, the closest contacts made by the copper atoms in the octahedral axial direction being in each case to atoms C(8) of adjacent molecules along [010], and of length 3.37 and 3.39 Å for molecules *A* and *B* respectively. The molecules are so oriented that the ethanolic hydroxyl groups of the independent molecules make contacts of 2.65 and 2.70 Å. The structure may then be described in terms of two-dimensional sheets of hydrogen-bonded molecules, parallel to (001). The compactness of this structure, resulting from the hydrogen bond formation, may be seen by comparing the density, 1.495 g.cm⁻³, with the values of 1.405 for bis-(*N*-ethylsalicylaldiminato)-copper (Clark, 1964) and 1.34 for bis-(*N*-butylsalicylaldiminato)copper (calculated from Frasson *et al.*, 1964).

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Refinement of the L-alanine crystal structure. By J.D. DUNITZ & R.R. RYAN, *Organic Chemistry Laboratory, Swiss Federal Institute of Technology, 8006 Zürich, Switzerland*

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Shortly after we had completed the measurement of a set of three-dimensional intensities for a crystal of L-alanine, we learned that the crystal structure had recently been determined by Simpson & Marsh (1966). In order to test the constancy of the molecular parameters derived from different data sources, we have carried out a series of full-matrix least-squares refinements with our data. We present here the comparison of our results with those of Simpson & Marsh (SM).

Our intensity measurements were made with a Hilger-Watts linear diffractometer, using Mo radiation with SrO/ZrO₂ balanced filters. The intensities of 522 independent reflexions were recorded in the layers *hk0*–*hk6* and converted to relative *F* values in the usual way. Absorption corrections were not deemed necessary.

Starting with SM's published parameters for the C, N and O atoms (hydrogen atoms were included in the structure factor calculations but not refined), our analysis leads to the results shown in Tables 1 and 2. The agreement is good as far as the chemical significance of the results is concerned; however, the differences, although small, seem statistically significant on the basis of the estimated standard deviations cited by SM. (We have not calculated the least-squares standard deviations of our parameters, but they should be of about the same order of magnitude as those of SM.) On the basis of tests using the function $R'' =$

We wish to thank Dr P.A. Vaughan of Rutgers University for his assistance with the refinement involving the anisotropic temperature factors.

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$[\sum w_i(F_o - F_c)^2 / \sum w_i F_o^2]^{1/2}$ (Hamilton, 1965) we find that our data reject the SM model at better than 0.005 level of significance, while the SM data reject our model at about the same level.

Table 1. *Positional parameters* ($\times 10^4$) *and bond lengths*

	<i>x</i> _{DR}	<i>x</i> _{SM}	σ_{SM}	<i>y</i> _{DR}	<i>y</i> _{SM}	σ_{SM}	<i>z</i> _{DR}	<i>z</i> _{SM}	σ_{SM}
O(1)	7278	7287	3	843	843	1	6280	6283	3
O(2)	4499	4501	3	1856	1850	1	7604	7609	3
C(1)	5606	5606	4	1413	1418	1	6023	6016	4
N	6565	6560	3	1375	1382	1	1853	1856	3
C(2)	4764	4769	4	1611	1612	1	3559	3563	4
C(3)	2744	2746	5	919	915	2	3021	3025	5

Bond	<i>d</i> _{DR}	<i>d</i> _{SM}
C(1)–O(1)	1.239 Å	1.247 Å
C(1)–O(2)	1.257	1.256
C(1)–C(2)	1.533	1.525
C(2)–C(3)	1.523	1.525
C(2)–N	1.496	1.491

$$\frac{R''(\text{SM data: our model})}{R''(\text{SM data: SM model})} = \frac{0.091}{0.070} = 1.30$$

$$\frac{R''(\text{our data: SM model})}{R''(\text{our data: our model})} = \frac{0.066}{0.049} = 1.34$$