

The Crystal and Molecular Structure of 9-Bromo-3,7,8,10-tetramethylisoalloxazine Monohydrate

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The crystal structure of 9-bromo-3,7,8,10-tetramethylisoalloxazine monohydrate, $C_{14}H_{13}BrN_4O_2 \cdot H_2O$, an example of a flavin derivative in an oxidized, neutral state, has been determined by three-dimensional X-ray analysis. The symmetry is monoclinic, space group $P2_1/c$, with unit cell constants $a = 13.047(5) \text{ \AA}$, $b = 7.053(5) \text{ \AA}$, $c = 17.957(5) \text{ \AA}$, and $\beta = 120.40(5)^\circ$. There are four molecules in the unit cell. The ring system within the molecule is fairly planar and the molecules are situated in layers normal to the b -axis. The structure was solved by the heavy-atom technique and the parameters obtained were refined by full matrix least squares analysis with a final reliability index, R , of 0.051 for the 1283 most significant observed reflections.

The structure analysis of 9-bromo-3,7,8,10-tetramethylisoalloxazine monohydrate was undertaken as part of a research program of X-ray structural studies of flavin compounds, important in many biological oxidation systems. A survey of the flavin compounds previously investigated within this project is given by Kierkegaard *et al.*¹ The present structure provides information on a flavin derivative in an oxidized nonprotonated state (Fig. 1). A brief description has appeared earlier.²

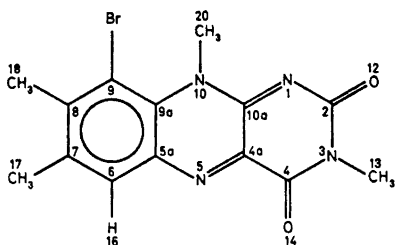


Fig. 1. Schematic drawing of 9-bromo-3,7,8,10-tetramethylisoalloxazine.

EXPERIMENTAL

Suitable single crystals were obtained by crystallization from an acetic acid solution of a specimen prepared by L. Maron³ at this Institute. A selected single crystal with the dimensions $0.3 \times 0.08 \times 0.02$ mm³ was indicated by Weissenberg methods to possess the monoclinic symmetry $P2_1/c$. The crystal data are given in Table 1. Three-dimensional

Table 1. Crystallographic data of 9-bromo-3,7,8,10-tetramethylisoalloxazine monohydrate. The estimated standard deviations are given in parentheses.

Lattice constants	$a = 13.047(5)$ Å $b = 7.053(5)$ Å $c = 17.957(5)$ Å $\beta = 120.40(5)^\circ$
Cell volume	$V = 1425.2$ Å ³
Density (X-ray)	$d = 1.7$ g cm ⁻³
Molecules per unit cell	$Z = 4$
Systematically absent reflections	$h0l$ when $l = 2n + 1$ $0k0$ when $k = 2n + 1$
Space group	$P2_1/c$
Wave length	$\lambda = 0.71069$ Å

X-ray intensity data were collected from a crystal mounted along the b -axis by a Siemens AED (Automatischer Einkristall-Diffraktometer) with Nb-filtered MoK radiation and a scintillation counter, with the pulse height analyzer set to collect about 90 % of the MoK α radiation. The $\theta - 2\theta$ scan technique was applied to yield integrated intensities. The intensities of 2906 reflections with $\theta \leq 25^\circ$ were determined as the averages of the intensities measured at two different symmetry related reciprocal lattice points. Only the 1283 most significant reflections were included in the further calculations. Lorentz, polarization and absorption corrections ($\mu = 30$ cm⁻¹)⁴ were applied to the net intensities.

STRUCTURE DETERMINATION AND REFINEMENT

The position of the bromine atom was determined from a three-dimensional Patterson synthesis and then refined with isotropic thermal parameter in a full matrix least squares refinement to an R -index

$$[R = \sum ||kF_o| - |F_c|| / \sum |kF_o|]$$

of 0.302. A three-dimensional difference electron density map displayed the atoms in the isoalloxazine ring near a plane $y = 0.6$. Successive three-dimensional difference electron density maps gave well-resolved peaks for the remaining nonhydrogen atoms including the oxygen atom of the water molecule. Full matrix least squares refinement of the atomic positions and the thermal parameters, anisotropic for the bromine atom and isotropic for the other nonhydrogen atoms, reduced the R -value to 0.077. After introduction of anisotropic temperature factors for all the 22 nonhydrogen atoms into the refinement, the R -index fell to 0.060.

Reasonable positions for all non-water hydrogen atoms were then obtained from a three-dimensional difference electron density map. Finally, three cycles of full matrix least squares refinement, with anisotropic temperature factors for all the 22 nonhydrogen atoms and constant isotropic (5 \AA^2) ones for the hydrogens, reduced the value of R to 0.051. Refinement was terminated when all shifts in the parameters were less than 5 % of their corresponding standard deviations. A subsequent difference electron density map did not

Table 2. Weight analysis obtained in the final cycle of the least squares refinement of 9-bromo-3,7,8,10-tetramethylisalloxazine monohydrate. The averages $\overline{w(|F_o| - |F_c|)^2} = \overline{w\Delta^2}$ are normalized. N is the number of independent reflections in each interval.

Interval $\sin \theta$	N	$\overline{w\Delta^2}$	Interval F_{obs}	N	$\overline{w\Delta^2}$
0.0 - 0.1973	233	0.9515	0.0 - 14.4	118	1.2247
0.1973 - 0.2485	202	1.0806	14.4 - 16.4	128	1.1049
0.2485 - 0.2845	175	1.1287	16.4 - 18.7	132	1.5215
0.2845 - 0.3131	156	0.7444	18.7 - 21.9	135	1.0459
0.3131 - 0.3373	158	0.8026	21.9 - 24.9	134	0.9258
0.3373 - 0.3585	110	0.8054	24.9 - 29.4	135	1.0770
0.3585 - 0.3774	98	1.1639	29.4 - 33.8	134	0.8165
0.3774 - 0.3945	83	1.2507	33.8 - 42.2	135	0.9311
0.3945 - 0.4103	62	1.3798	42.2 - 57.8	134	0.6263
0.4103 - 0.4250	43	1.1055	57.8 - 304.0	135	0.7671

Table 3. Final positional parameters for the nonhydrogen atoms with estimated standard deviations.

Atom	x	y	z
Br	0.7777(1)	0.5548(2)	0.49550(6)
N(1)	0.3090(7)	0.6316(14)	0.3429(5)
C(2)	0.1872(9)	0.6219(16)	0.2874(6)
N(3)	0.1377(6)	0.6024(13)	0.1989(5)
C(4)	0.2021(8)	0.6053(14)	0.1575(6)
C(4A)	0.3328(7)	0.6183(14)	0.2165(5)
N(5)	0.3987(7)	0.6280(13)	0.1817(4)
C(5A)	0.5203(7)	0.6386(15)	0.2363(6)
C(6)	0.5908(7)	0.6364(13)	0.1978(5)
C(7)	0.7123(8)	0.6363(15)	0.2467(6)
C(8)	0.7668(8)	0.6182(15)	0.3374(6)
C(9)	0.6977(8)	0.6154(14)	0.3776(6)
C(9A)	0.5728(8)	0.6363(15)	0.3274(6)
N(10)	0.4964(6)	0.6526(12)	0.3618(4)
C(10A)	0.3771(8)	0.6303(16)	0.3078(6)
O(12)	0.1219(6)	0.6241(13)	0.3179(5)
C(13)	0.0076(8)	0.5863(19)	0.1444(6)
O(14)	0.1590(5)	0.5907(12)	0.0801(4)
C(17)	0.7833(8)	0.6349(17)	0.2018(6)
C(18)	0.9010(8)	0.6035(18)	0.3922(6)
C(20)	0.5384(9)	0.7216(19)	0.4504(6)
O	0.2989(11)	0.6022(15)	-0.0173(6)

Table 4. Anisotropic thermal parameters for the nonhydrogen atoms and their standard deviations. ($T = \exp[-(\beta_{11}h^2 + \beta_{22}k^2 + \beta_{33}l^2 + \beta_{12}hk + \beta_{13}hl + \beta_{23}kl)]$.)

Atom	β_{11}	β_{22}	β_{33}	β_{12}	β_{13}	β_{23}
Br	0.00583(9)	0.0285(3)	0.00235(4)	-0.00207(28)	0.00159(9)	-0.00241(26)
N(1)	0.0054(7)	0.0270(26)	0.0028(3)	0.0021(21)	0.0045(8)	0.0001(15)
C(2)	0.0061(8)	0.0206(29)	0.0032(4)	0.0052(24)	0.0060(10)	0.0025(17)
N(3)	0.0037(6)	0.0269(29)	0.0033(3)	0.0020(19)	0.0042(8)	0.0035(15)
C(4)	0.0045(7)	0.0155(31)	0.0027(4)	0.0023(21)	0.0035(9)	0.0022(14)
C(4A)	0.0035(7)	0.0153(25)	0.0027(4)	-0.0013(20)	0.0036(8)	-0.0002(15)
N(5)	0.0052(6)	0.0220(23)	0.0015(3)	-0.0004(19)	0.0018(7)	0.0003(12)
C(5A)	0.0024(7)	0.0165(23)	0.0027(4)	0.0018(20)	0.0016(8)	0.0015(15)
C(6)	0.0059(7)	0.0194(21)	0.0020(3)	0.0021(19)	0.0031(7)	-0.0004(13)
C(7)	0.0048(8)	0.0148(24)	0.0037(5)	0.0006(22)	0.0040(10)	0.0030(17)
C(8)	0.0052(8)	0.0140(27)	0.0035(4)	-0.0001(22)	0.0035(10)	-0.0015(16)
C(9)	0.0045(7)	0.0162(26)	0.0029(4)	-0.0016(21)	0.0029(9)	0.0029(16)
C(9A)	0.0048(7)	0.0118(22)	0.0027(4)	-0.0020(21)	0.0039(9)	-0.0001(15)
N(10)	0.0048(6)	0.0173(20)	0.0024(3)	0.0008(19)	0.0036(7)	0.0014(13)
C(10A)	0.0049(8)	0.0198(26)	0.0024(4)	-0.0007(23)	0.0040(9)	-0.0002(15)
O(12)	0.0070(6)	0.0356(26)	0.0051(4)	0.0023(20)	0.0090(9)	0.0005(16)
C(13)	0.0049(7)	0.0254(33)	0.0041(4)	-0.0027(24)	0.0038(8)	-0.0019(19)
O(14)	0.0050(5)	0.0403(29)	0.0020(3)	-0.0072(20)	0.0023(6)	-0.0026(13)
C(17)	0.0062(8)	0.0262(27)	0.0040(4)	-0.0055(24)	0.0070(10)	-0.0054(18)
C(18)	0.0044(6)	0.0352(40)	0.0038(4)	-0.0047(25)	0.0028(7)	-0.0030(20)
C(20)	0.0074(9)	0.0290(36)	0.0035(4)	-0.0041(28)	0.0044(10)	-0.0052(19)
O	0.0244(16)	0.0380(35)	0.0072(6)	0.0182(36)	0.0186(17)	0.0039(22)

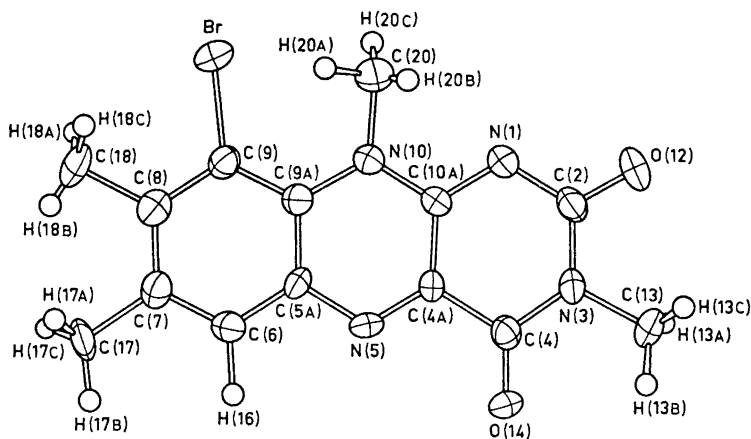


Fig. 2. Molecular conformation observed in the crystal structure of 9-bromo-3,7,8,10-tetramethylisoalloxazine monohydrate with atoms numbered for reference in the text.

reveal the positions of the two remaining hydrogens (bonded to the water oxygen).

The scattering factors for bromine, carbon, nitrogen, and oxygen were taken from Hanson *et al.*⁵ and that for hydrogen were given by Stewart *et al.*⁶

Table 5. Atomic coordinates for hydrogen atoms with their standard deviations and isotropic thermal parameters (\AA^2).

Atom	<i>x</i>	<i>y</i>	<i>z</i>	<i>B</i>
H(16)	0.544(10)	0.621(17)	0.124(8)	5.00
H(13A)	-0.008(9)	0.488(16)	0.149(7)	5.00
H(13B)	-0.022(8)	0.681(14)	0.083(6)	5.00
H(13C)	-0.027(8)	0.706(15)	0.161(6)	5.00
H(17A)	0.835(9)	0.733(15)	0.226(6)	5.00
H(17B)	0.738(8)	0.619(13)	0.135(6)	5.00
H(17C)	0.844(9)	0.499(13)	0.221(6)	5.00
H(18A)	0.926(8)	0.447(14)	0.431(6)	5.00
H(18B)	0.913(9)	0.602(15)	0.353(6)	5.00
H(18C)	0.914(8)	0.705(15)	0.434(6)	5.00
H(20A)	0.613(8)	0.823(15)	0.464(6)	5.00
H(20B)	0.489(9)	0.799(15)	0.441(7)	5.00
H(20C)	0.563(8)	0.594(15)	0.489(6)	5.00

Table 6. Intramolecular bond distances (in \AA) and their estimated standard deviations.

Br - C(9)	1.876(9)	N(3) - C(13)	1.473(11)
N(1) - C(2)	1.385(12)	C(4) - O(14)	1.212(10)
C(2) - N(3)	1.387(12)	C(7) - C(17)	1.505(13)
N(3) - C(4)	1.376(11)	C(8) - C(18)	1.516(13)
C(4) - C(4A)	1.487(12)	N(10) - C(20)	1.479(12)
C(4A) - N(5)	1.295(11)	C(6) - H(16)	1.14(12)
N(5) - C(5A)	1.382(11)	C(13) - H(13A)	0.74(11)
C(5A) - C(6)	1.403(11)	C(13) - H(13B)	1.18(10)
C(6) - C(7)	1.369(12)	C(13) - H(13C)	1.07(10)
C(7) - C(8)	1.415(13)	C(17) - H(17A)	0.91(10)
C(8) - C(9)	1.411(13)	C(17) - H(17B)	1.04(10)
C(9) - C(9A)	1.416(13)	C(17) - H(17C)	1.18(10)
C(9A) - N(10)	1.418(11)	C(18) - H(18A)	1.25(10)
N(10) - C(10A)	1.363(12)	C(18) - H(18B)	0.80(10)
C(10A) - N(1)	1.325(12)	C(18) - H(18C)	0.98(10)
C(4A) - C(10A)	1.438(12)	C(20) - H(20A)	1.13(10)
C(5A) - C(9A)	1.418(12)	C(20) - H(20B)	0.79(11)
C(2) - O(12)	1.224(12)	C(20) - H(20C)	1.08(10)

Table 7. Intramolecular bond angles (in degrees) with their estimated standard deviations.

Br - C(9) - C(9A)	123.1(7) $^\circ$	C(6) - C(7) - C(8)	118.9(8)
Br - C(9) - C(8)	116.5(7)	C(6) - C(7) - C(17)	118.9(8)
N(1) - C(2) - N(3)	122.0(8)	C(7) - C(8) - C(9)	120.7(8)
N(1) - C(2) - O(12)	118.8(8)	C(7) - C(8) - C(18)	119.8(8)
C(2) - N(3) - C(13)	118.7(7)	C(8) - C(9) - C(9A)	120.0(8)
C(2) - N(3) - C(4)	124.0(7)	C(9) - C(9A) - N(10)	124.6(8)
N(3) - C(4) - O(14)	124.2(8)	C(9A) - N(10) - C(20)	122.4(7)
N(3) - C(4) - C(4A)	114.2(7)	C(9A) - N(10) - C(10A)	118.8(7)
C(4) - C(4A) - C(10A)	118.1(7)	C(9A) - C(5A) - C(6)	120.9(7)
C(4) - C(4A) - N(5)	117.4(8)	N(10) - C(10A) - N(1)	117.4(8)
C(4A) - C(10A) - N(1)	124.3(8)	C(10A) - N(1) - C(2)	117.3(8)
C(4A) - C(10A) - N(10)	118.2(8)	C(10A) - C(4A) - N(5)	124.4(8)
C(4A) - N(5) - C(5A)	117.8(7)	O(12) - C(2) - N(3)	119.2(9)
N(5) - C(5A) - C(9A)	121.8(7)	C(13) - N(3) - C(4)	117.2(7)
N(5) - C(5A) - C(6)	117.0(7)	O(14) - C(4) - C(4A)	121.5(8)
C(5A) - C(9A) - N(10)	117.8(7)	C(17) - C(7) - C(8)	121.9(8)
C(5A) - C(9A) - C(9)	117.6(8)	C(18) - C(8) - C(9)	119.5(8)
C(5A) - C(6) - C(7)	121.2(7)	C(20) - N(10) - C(10A)	118.1(7)

Table 9. Computer programs used for the crystallographic calculations.

Program name and function	Authors
PIRUM. Indexing of powder photographs and least squares refinement of unit cell parameters.	Werner, P.-E.
SIP. Generation of steering paper tape for the SIEMENS AED.	Norrestam, R.
SIMS. Interpretation of output on paper tape from the SIEMENS AED and evaluation of intensities.	Norrestam, R.
DATA P 2. Lp- and absorption corrections. Preparative calculations for extinction correction.	Coppens P., Leiserowitz, L. & Rabinovich, D. Modified by Olofsson, O., Elfström, M., Brandt, B., Åsbrink, S. & Nord, A.
DATA. Reflection data handling.	Brandt, B.
DRF. Fourier summations and structure factor calculations.	Zalkin, A. Modified by Liminga, R., Lundgren, J.-O., Lindgren, O., Brandt, B. & Nord, A.
LALS. Full matrix least squares refinement of positional and thermal parameters and of scale factors.	Gantzel, P. K., Sparks, R. A. & Trueblood, K. N. Modified by Zalkin, A., Lundgren, J.-O., Liminga, R., Brändén, C.-I., Lindgren, O., Brandt, B. & Nord, A.
DISTAN. Calculation of interatomic distances, and bond angles with estimated standard deviations.	Zalkin, A. Modified by Nord, A. & Brandt, B.
INERT. Calculation of least squares plane and axes of inertia.	Norrestam, R.
ORTEP. Thermal-ellipsoid plot for crystal structure illustrations.	Johnson, C. K. Modified by Carlbon, I.
ACTA. Editing of structure factor tables.	Norrestam, R.
SCF-OPSZDO. Self-consistent field molecular orbital calculations.	Sundbom, M. & Roos, B.

Hughes' ⁷ weighting scheme was used in all least squares refinements, with $|F_{o,\min}| = 10.0$ and $h = 4.0$. The weight analysis obtained in the last cycle is given in Table 2. The positional and thermal parameters (with estimated standard deviations, e.s.d.'s) are given in Tables 3, 4, and 5, intramolecular distances and angles in Tables 6 and 7, and structure factors in Table 8. The atomic labeling is shown in Fig. 2. The computer programs are identified in Table 9; most of the calculations were performed on an IBM-360/75 computer.

DESCRIPTION AND DISCUSSION OF THE STRUCTURE

The molecular packing viewed along the b and c axes is shown in Figs. 3a and 3b. The crystal structure is built up of an infinite stacking in the $[010]$ direction of almost planar 9-bromo-3,7,8,10-tetramethylisalloxazine molecules with parallel molecular orientation. The intermolecular distance in the $[010]$ direction is about 3.5 Å, a normal van der Waals separation (usual value 3.5–3.7 Å).

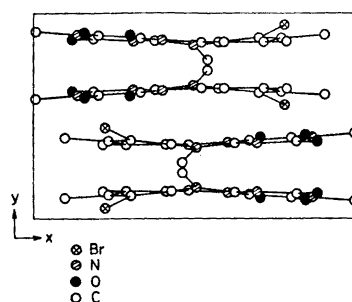
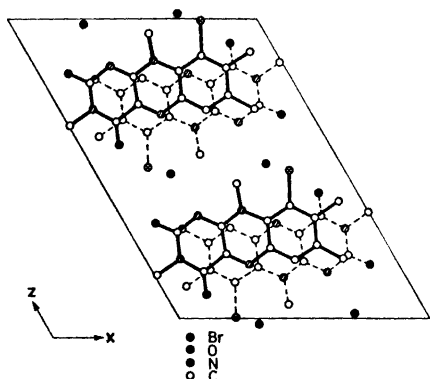


Fig. 3a. A packing diagram, excluding hydrogen atoms, viewed along the b axis.

Fig. 3b. A packing diagram, excluding hydrogen atoms, viewed along the c axis.

Table 10. Intermolecular distances shorter than 3.6 Å between nonhydrogen atoms and their estimated standard deviations.

Code for symmetry related atoms		Atom at		Superscript		Atom at		
Superscript	x	y	z	Superscript	x	$\frac{1}{2}-y$	$\frac{1}{2}+z$	
None	x	y	z	iii	x	$\frac{1}{2}-y$	$\frac{1}{2}+z$	
i	$1-x$	$1-y$	$1-z$	iv	$1-x$	$\frac{1}{2}+y$	$\frac{1}{2}-z$	
ii	$1-x$	$-\frac{1}{2}+y$	$\frac{1}{2}-z$	v	$-1+x$	y	z	
				vi	$1-x$	$1-y$	$-z$	
	Br	-O(12 i)			3.177(8) Å			
	Br	-O(ii)			3.427(11)			
	N(1)	-O(iii)			3.189(13)			
	C(2)	-C(17 ii)			3.451(16)			
	C(4)	-C(8 ii)			3.455(14)			
	C(4A)	-C(7 ii)			3.569(15)			
	N(5)	-C(9A ii)			3.500(14)			
	N(5)	-O			3.130(12)			
	C(6)	-N(10 ii)			3.585(13)			
	C(6)	-C(10A iv)			3.516(15)			
	C(8)	-O(14 iv)			3.576(13)			
	O(12)	-O(iii)			3.301(13)			
	C(13)	-C(17 v)			3.577(13)			
	O(14)	-O			3.103(13)			
	C(17)	-O(vi)			3.364(14)			
	C(20)	-O(iv)			3.251(17)			

The unit cell contains two molecular stacks related by inversions. Each stack is built up by 9-bromo-3,7,8,10-tetramethylisoalloxazine molecules with alternating orientation, as seen in Fig. 3a. The water molecules, located between the stacks, seem to be involved in several weak hydrogen bonds, linking the flavin molecules together in the [001] direction. Since the positions of the water hydrogens could not be determined it is not possible to draw any firm conclusions regarding the hydrogen bond scheme. The water oxygen forms several contacts less than 3.6 Å (*cf.* Table 10) *viz.* Br—O(ii), N(1)—O(iii), N(5)—O, O(12)—O(iii), O(14)—O, C(17)—O(vi), and C(20)—O(iv). Of these, the first five ranging from 3.10 to 3.43 Å, might be caused by hydrogen bonds. Since the 9-bromo-3,7,8,10-tetramethylisoalloxazine molecule has no suitable donors to hydrogen bonds, the water molecule might serve as a donor to weak hydrogen bonds. Judging from the distances 3.13 and 3.10 Å, respectively, it is likely that the water molecule forms hydrogen bonds (possibly bifurcated) at least to the atoms N(5) and O(14) of the flavin molecule. It is interesting to note that this position of the water molecule relative to the flavin nucleus is the same as that suggested for metal ions in metal flavin complexes from spectroscopical investigations (see, *e.g.*, Hemmerich and Spence⁸). The lack of strong intermolecular contacts of the water molecule is probably the cause of the large temperature vibrations of the water oxygen and the problems of locating its hydrogens.

Apart from the contacts in the [001] direction involving the water molecule given above, there is one remarkably short intermolecular distance, 3.18 Å, between a bromine of one flavin molecule and a keto oxygen, O(12), of another. The occurrence of this short contact might be the reason that no strong hydrogen bonds are formed in the crystal structure, *viz.* if the observed packing is energetically favourable as compared with different packings which allow stronger hydrogen bonds. The shortest intermolecular distance in the [100] direction is 3.58 Å, a reasonable van der Waals contact in this direction.

As seen in Fig. 3b the isoalloxazine ring system of the flavin molecule is almost planar and the major deviations from planarity is obtained for the atoms C(20) and Br. Least squares planes calculated through the individual rings of the flavin molecule are given in Table 11. The angle between the normals of the benzenoid and pyrazinoid rings is 7.0°, and the angle between the normals of the pyrazinoid and pyrimidinoid rings is 1.0°. The angle between the planes of the benzenoid and pyrazinoid rings arises from a slight twist of the isoalloxazine ring, attributable to repulsive forces between the bromine atom and the methyl group attached to N(10). Thus the Br—C(20) distance becomes 3.04 Å, equal to the Br—C(18) distance of 3.03 Å. The conclusion that the observed distortion of the isoalloxazine ring in the present study is caused by overcrowding effects (due to the 9-bromo-substitution of the benzenoid ring) is supported by details of the crystal structure of 3-methyl-lumiflavin⁹ determined at this Institute.

In Table 12 the intramolecular bond distances found in the present study are compared with those obtained from self-consistent field molecular orbital calculations of the Pariser-Parr-Pople (PPP) type. The molecular orbitals have been evaluated by means of a computer program kindly put at our disposal by Marianne Sundbom (Institute of Theoretical Physics, University of Stock-

Table 11. Deviations of the atoms from least squares planes. The planes are defined by $Ax + By + Cz = D$, where x, y, z are in Å units along the axes a^* , b and c . The atoms indicated with asterisks were omitted from the calculations of the least squares planes.

Plane I	Atom	Deviation	Plane III	Atom	Deviation
A = -0.0235	N(1)	0.038 Å	A = -0.0546	C(4A)	-0.032 Å
B = 0.9996	C(2)	0.005	B = 0.9981	N(5)	0.027
C = -0.0156	N(3)	-0.100	C = -0.0302	C(5A)	0.022
D = 4.2691 Å	C(4)	-0.078	D = 4.1291 Å	C(9A)	-0.065
	C(4A)	-0.024		N(10)	0.062
	N(5)	0.043		C(10A)	-0.015
	C(5A)	0.083		N(1)*	0.003
	C(6)	0.067		C(4)*	-0.037
	C(7)	0.033		C(6)*	-0.002
	C(8)	-0.129		C(9)*	-0.292
	C(9)	-0.149		C(20)*	0.483
	C(9A)	0.033			
	N(10)	0.150			
	C(10A)	0.028			
	O(12)*	0.022			
	C(13)*	-0.177			
	O(14)*	-0.152			
	C(17)*	0.024			
	C(18)*	-0.270			
	C(20)*	0.606			
	Br*	-0.622			
	O*	-0.067			
Plane II	Atom	Deviation	Plane IV	Atom	Deviation
A = -0.0603	N(1)	-0.004 Å	A = 0.0504	C(5A)	-0.014 Å
B = 0.9970	C(2)	0.019	B = 0.9982	C(6)	-0.029
C = -0.0483	N(3)	-0.023	C = 0.0339	C(7)	0.042
D = 4.0370 Å	C(4)	0.010	D = 4.8325 Å	C(8)	-0.011
	C(4A)	0.004		C(9)	-0.031
	C(10A)	-0.007		C(9A)	0.043
	O(12)*	0.032		N(5)*	-0.164
	C(13)*	-0.042		N(10)*	0.152
	O(14)*	-0.010		C(17)*	0.029
	N(5)*	0.078		C(18)*	-0.036
	N(10)*	0.060		Br*	-0.358

holm). The input atomic coordinates used were derived from those obtained in the present study by assuming a strictly planar conformation of the molecule. The other input parameters used were chosen in accordance with those suggested by Roos and Skancke,¹⁰ Roos,¹¹ Fischer-Hjalmars and Sundbom,¹² and Jensen and Skancke.¹³ The bond lengths were derived from bond order-bond length correlations, *viz.*

$$R_{CC} = 1.517 - 0.180 p_{CC}$$

$$R_{CN} = 1.458 - 0.180 p_{CN}$$

$$R_{CO} = 1.365 - 0.180 p_{CO}$$

Table 12. Calculated π -bond orders with corresponding bond lengths compared with those obtained in the present study.

	π -b. o.	d_{calc}	d_{obs}
N(1) – C(2)	0.42	1.383 Å	1.385 Å
C(2) – N(3)	0.44	1.380	1.387
N(3) – C(4)	0.50	1.368	1.376
C(4) – C(4A)	0.27	1.468	1.487
C(4A) – N(5)	0.82	1.311	1.295
N(5) – C(5A)	0.44	1.379	1.382
C(5A) – C(6)	0.57	1.415	1.403
C(6) – C(7)	0.72	1.388	1.371
C(7) – C(8)	0.60	1.409	1.415
C(8) – C(9)	0.65	1.399	1.411
C(9) – C(9A)	0.58	1.413	1.416
C(9A) – C(5A)	0.59	1.411	1.418
C(9A) – N(10)	0.37	1.391	1.418
N(10) – C(10A)	0.41	1.384	1.363
C(4A) – C(10A)	0.36	1.453	1.438
C(10A) – N(1)	0.77	1.319	1.325
C(2) – O(12)	0.68	1.243	1.224
N(3) – C(13)	0.05	1.449	1.473
C(4) – O(14)	0.68	1.242	1.212
C(7) – C(17)	0.14	1.492	1.506
C(8) – C(18)	0.17	1.486	1.516
N(10) – C(20)	0.05	1.449	1.479

where p is the calculated π -bond order and R the corresponding bond length.

The agreement between the distances found in the crystal structure and those obtained from the molecular orbital calculations is encouraging; the differences among the endocyclic bonds are all less than 0.03 Å, well below three times the estimated standard deviations of the corresponding bond lengths. The main features of the evaluated π -bond order scheme are in accordance with the commonly accepted bonding scheme for an oxidized flavin derivative as given in Fig. 1. Thus the highest π -bond orders within the ring system are those between C(4a) – N(5) (0.82) and N(1) – C(10A) (0.77). It is obvious that both oxygens O(12) and O(14) attached to the pyrimidinoid ring are of keto type.

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