

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 $a = 13.047(5)$ Å, $b = 7.053(5)$ Å, $c = 17.957(5)$ Å, 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

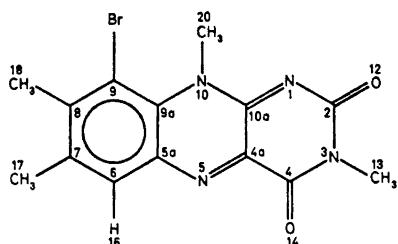


Fig. 1. Schematic drawing of 9-bromo-3,7,8,10-tetramethylisoalloxazine.

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-tetramethylisoalloxazine monohydrate. The averages $w(|F_o| - |F_c|)^2 = w\bar{A}^2$ are normalized. N is the number of independent reflections in each interval.

Interval $\sin \theta$	N	$\overline{w\bar{A}^2}$	Interval F_{obs}	N	$\overline{w\bar{A}^2}$
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)

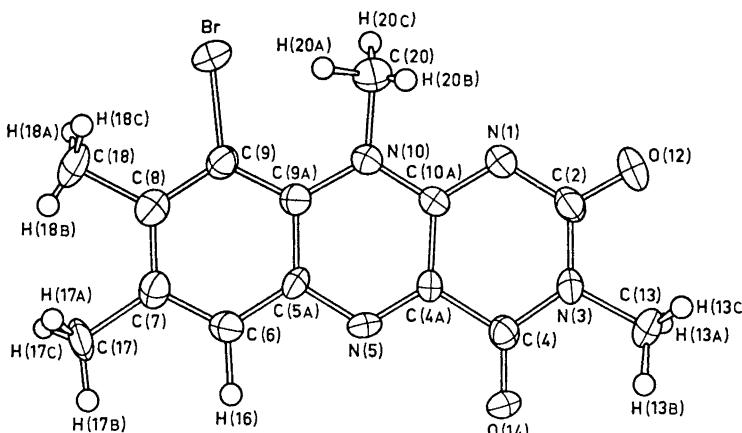


Fig. 2. Molecular conformation observed in the crystal structure of 9-bromo-3,7,8,10-tetramethylisoalloxazine monohydrate with atoms numbered for reference in the text.

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) \AA	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 8. Structure factors. Each of three columns contains k , $10|F_o|$, and $10|F_c|$, and is headed by the values of h and l common to the group.

h	0	$L = 0$	a	$53H$	572	1	259	261	5	194	205	7	139	114	5	198	223	$H = 13$	$L = 0$	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45	46	47	48	49	50	51	52	53	54	55	56	57	58	59	60	61	62	63	64	65	66	67	68	69	70	71	72	73	74	75	76	77	78	79	80	81	82	83	84	85	86	87	88	89	90	91	92	93	94	95	96	97	98	99	100	101	102	103	104	105	106	107	108	109	110	111	112	113	114	115	116	117	118	119	120	121	122	123	124	125	126	127	128	129	130	131	132	133	134	135	136	137	138	139	140	141	142	143	144	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160	161	162	163	164	165	166	167	168	169	170	171	172	173	174	175	176	177	178	179	180	181	182	183	184	185	186	187	188	189	190	191	192	193	194	195	196	197	198	199	200	201	202	203	204	205	206	207	208	209	210	211	212	213	214	215	216	217	218	219	220	221	222	223	224	225	226	227	228	229	230	231	232	233	234	235	236	237	238	239	240	241	242	243	244	245	246	247	248	249	250	251	252	253	254	255	256	257	258	259	260	261	262	263	264	265	266	267	268	269	270	271	272	273	274	275	276	277	278	279	280	281	282	283	284	285	286	287	288	289	290	291	292	293	294	295	296	297	298	299	300	301	302	303	304	305	306	307	308	309	310	311	312	313	314	315	316	317	318	319	320	321	322	323	324	325	326	327	328	329	330	331	332	333	334	335	336	337	338	339	340	341	342	343	344	345	346	347	348	349	350	351	352	353	354	355	356	357	358	359	360	361	362	363	364	365	366	367	368	369	370	371	372	373	374	375	376	377	378	379	380	381	382	383	384	385	386	387	388	389	390	391	392	393	394	395	396	397	398	399	400	401	402	403	404	405	406	407	408	409	410	411	412	413	414	415	416	417	418	419	420	421	422	423	424	425	426	427	428	429	430	431	432	433	434	435	436	437	438	439	440	441	442	443	444	445	446	447	448	449	450	451	452	453	454	455	456	457	458	459	460	461	462	463	464	465	466	467	468	469	470	471	472	473	474	475	476	477	478	479	480	481	482	483	484	485	486	487	488	489	490	491	492	493	494	495	496	497	498	499	500	501	502	503	504	505	506	507	508	509	510	511	512	513	514	515	516	517	518	519	520	521	522	523	524	525	526	527	528	529	530	531	532	533	534	535	536	537	538	539	540	541	542	543	544	545	546	547	548	549	550	551	552	553	554	555	556	557	558	559	560	561	562	563	564	565	566	567	568	569	570	571	572	573	574	575	576	577	578	579	580	581	582	583	584	585	586	587	588	589	590	591	592	593	594	595	596	597	598	599	600	601	602	603	604	605	606	607	608	609	610	611	612	613	614	615	616	617	618	619	620	621	622	623	624	625	626	627	628	629	630	631	632	633	634	635	636	637	638	639	640	641	642	643	644	645	646	647	648	649	650	651	652	653	654	655	656	657	658	659	660	661	662	663	664	665	666	667	668	669	670	671	672	673	674	675	676	677	678	679	680	681	682	683	684	685	686	687	688	689	690	691	692	693	694	695	696	697	698	699	700	701	702	703	704	705	706	707	708	709	710	711	712	713	714	715	716	717	718	719	720	721	722	723	724	725	726	727	728	729	730	731	732	733	734	735	736	737	738	739	740	741	742	743	744	745	746	747	748	749	750	751	752	753	754	755	756	757	758	759	760	761	762	763	764	765	766	767	768	769	770	771	772	773	774	775	776	777	778	779	780	781	782	783	784	785	786	787	788	789	790	791	792	793	794	795	796	797	798	799	800	801	802	803	804	805	806	807	808	809	810	811	812	813	814	815	816	817	818	819	820	821	822	823	824	825	826	827	828	829	830	831	832	833	834	835	836	837	838	839	840	841	842	843	844	845	846	847	848	849	850	851	852	853	854	855	856	857	858	859	860	861	862	863	864	865	866	867	868	869	870	871	872	873	874	875	876	877	878	879	880	881	882	883	884	885	886	887	888	889	890	891	892	893	894	895	896	897	898	899	900	901	902	903	904	905	906	907	908	909	910	911	912	913	914	915	916	917	918	919	920	921	922	923	924	925	926	927	928	929	930	931	932	933	934	935	936	937	938	939	940	941	942	943	944	945	946	947	948	949	950	951	952	953	954	955	956	957	958	959	960	961	962	963	964	965	966	967	968	969	970	971	972	973	974	975	976	977	978	979	980	981	982	983	984	985	986	987	988	989	990	991	992	993	994	995	996	997	998	999	1000	1001	1002	1003	1004	<

Table 8. Continued.

H	145	11	H	+	L=7	H	295	L=11	H	6	L=14	H	7	L=19	H	9	L=6	H	10	L=13	H	220	204	
H	265	25	H	+	L=7	H	295	L=11	H	6	L=14	H	7	L=19	H	9	L=6	H	10	L=13	H	200	217	
H	1093	145	H	+	L=7	H	426	L=7	H	6	L=14	H	7	L=19	H	9	L=6	H	10	L=13	H	12	L=7	
H	383	345	H	+	L=7	H	342	L=7	H	6	L=14	H	7	L=19	H	9	L=6	H	10	L=13	H	230	214	
H	145	145	H	+	L=7	H	6	L=14	H	6	L=15	H	7	L=19	H	9	L=7	H	10	L=13	H	12	L=8	
H	145	15	H	+	L=7	H	6	L=14	H	6	L=15	H	7	L=19	H	9	L=7	H	10	L=13	H	12	L=8	
H	145	213	H	+	L=7	H	6	L=14	H	6	L=15	H	7	L=19	H	9	L=7	H	10	L=13	H	3	238	264
H	164	134	H	+	L=7	H	6	L=14	H	6	L=15	H	7	L=19	H	9	L=7	H	10	L=13	H	1	137	41
H	9	241	H	+	L=7	H	195	L=16	H	6	L=14	H	7	L=19	H	9	L=7	H	10	L=13	H	1	134	286
H	32	L=11	H	+	L=7	H	342	L=12	H	6	L=14	H	7	L=19	H	9	L=7	H	10	L=13	H	1	137	41
H	221	510	H	+	L=7	H	345	L=14	H	6	L=14	H	7	L=19	H	9	L=7	H	10	L=13	H	12	L=12	
H	271	251	H	+	L=7	H	200	L=17	H	548	L=19	H	7	L=20	H	9	L=8	H	10	L=17	H	12	L=10	
H	372	369	H	+	L=7	H	367	L=19	H	402	L=20	H	7	L=20	H	9	L=8	H	10	L=17	H	12	L=10	
H	141	172	H	+	L=7	H	121	L=14	H	168	L=14	H	6	L=17	H	2	L=21	H	10	L=18	H	12	L=11	
H	10	L=10	H	+	L=7	H	294	L=11	H	359	L=13	H	6	L=18	H	2	L=20	H	10	L=19	H	12	L=10	
H	10	524	H	+	L=7	H	308	L=12	H	308	L=16	H	6	L=18	H	2	L=20	H	10	L=19	H	12	L=12	
H	235	251	H	+	L=7	H	196	L=16	H	300	L=16	H	6	L=18	H	2	L=20	H	10	L=19	H	12	L=12	
H	203	149	H	+	L=7	H	196	L=21	H	300	L=20	H	6	L=18	H	2	L=20	H	10	L=19	H	12	L=12	
H	317	L=11	H	+	L=7	H	329	L=28	H	367	L=11	H	6	L=19	H	2	L=21	H	10	L=19	H	12	L=12	
H	216	200	H	+	L=7	H	326	L=26	H	367	L=24	H	6	L=19	H	2	L=21	H	10	L=19	H	12	L=12	
H	206	242	H	+	L=7	H	274	L=29	H	367	L=24	H	6	L=19	H	2	L=21	H	10	L=19	H	12	L=12	
H	378	L=12	H	+	L=7	H	329	L=16	H	307	L=16	H	6	L=19	H	2	L=21	H	10	L=19	H	12	L=12	
H	257	249	H	+	L=7	H	329	L=10	H	305	L=16	H	6	L=19	H	2	L=21	H	10	L=19	H	12	L=12	
H	263	271	H	+	L=7	H	307	L=34	H	210	L=21	H	7	L=22	H	3	L=29	H	11	L=31	H	12	L=31	
H	143	73	H	+	L=7	H	278	L=27	H	340	L=24	H	6	L=20	H	2	L=21	H	10	L=19	H	12	L=12	
H	316	L=13	H	+	L=7	H	249	L=17	H	326	L=20	H	6	L=20	H	2	L=21	H	10	L=19	H	12	L=12	
H	209	202	H	+	L=7	H	316	L=22	H	316	L=19	H	6	L=20	H	2	L=21	H	10	L=19	H	12	L=12	
H	144	134	H	+	L=7	H	338	L=24	H	362	L=16	H	6	L=20	H	2	L=21	H	10	L=19	H	12	L=12	
H	333	L=14	H	+	L=7	H	150	L=17	H	328	L=24	H	6	L=20	H	2	L=21	H	10	L=19	H	12	L=12	
H	242	252	H	+	L=7	H	150	L=17	H	328	L=24	H	6	L=20	H	2	L=21	H	10	L=19	H	12	L=12	
H	193	193	H	+	L=7	H	205	L=20	H	368	L=20	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	241	262	H	+	L=7	H	150	L=19	H	338	L=20	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	169	190	H	+	L=7	H	277	L=21	H	368	L=20	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	370	365	H	+	L=7	H	167	L=18	H	329	L=21	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	205	203	H	+	L=7	H	1009	L=76	H	329	L=30	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	405	414	H	+	L=7	H	129	L=23	H	302	L=26	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	224	226	H	+	L=7	H	454	L=21	H	323	L=24	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	214	168	H	+	L=7	H	494	L=21	H	323	L=24	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	234	234	H	+	L=7	H	369	L=31	H	369	L=32	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	3	205	H	+	L=7	H	305	L=28	H	368	L=28	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	241	262	H	+	L=7	H	150	L=19	H	328	L=24	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	370	L=16	H	+	L=7	H	167	L=18	H	329	L=21	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	225	286	H	+	L=7	H	1009	L=76	H	329	L=26	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	1	405	L=17	H	+	L=7	H	129	L=23	H	302	L=26	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12
H	204	226	H	+	L=7	H	504	L=20	H	323	L=24	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	242	254	H	+	L=7	H	494	L=21	H	323	L=24	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	1	234	L=19	H	+	L=7	H	255	L=27	H	368	L=30	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12
H	3	210	L=20	H	+	L=7	H	105	L=17	H	329	L=26	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12
H	4	754	L=21	H	+	L=7	H	150	L=21	H	368	L=24	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12
H	229	234	H	+	L=7	H	208	L=21	H	368	L=24	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	235	256	H	+	L=7	H	493	L=21	H	368	L=24	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	148	149	H	+	L=7	H	359	L=38	H	180	L=13	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	761	750	H	+	L=7	H	258	L=29	H	103	L=100	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	362	364	H	+	L=7	H	100	L=13	H	376	L=84	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	239	254	H	+	L=7	H	1013	L=46	H	336	L=92	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	1070	1042	H	+	L=7	H	439	L=45	H	747	L=705	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	236	222	H	+	L=7	H	144	L=39	H	479	L=486	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	279	292	H	+	L=7	H	146	L=44	H	337	L=97	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	200	190	H	+	L=7	H	140	L=44	H	140	L=97	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	5	L=6	H	+	L=7	H	1075	L=46	H	6	L=9	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	428	429	H	+	L=7	H	1075	L=46	H	6	L=9	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	262	265	H	+	L=7	H	314	L=20	H	6	L=10	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	204	205	H	+	L=7	H	314	L=20	H	6	L=10	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	715	716	H	+	L=7	H	302	L=13	H	6	L=10	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	240	241	H	+	L=7	H	302	L=13	H	6	L=10	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	226	246	H	+	L=7	H	302	L=13	H	6	L=10	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	146	168	H	+	L=7	H	511	L=95	H	6	L=11	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	269	270	H	+	L=7	H	369	L=95	H	6	L=11	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	205	210	H	+	L=7	H	369	L=95	H	6	L=11	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	428	429	H	+	L=7	H	138	L=80	H	6	L=11	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	262	265	H	+	L=7	H	314	L=20	H	6	L=11	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	204	205	H	+	L=7	H	314	L=20	H	6	L=11	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	715	716	H	+	L=7	H	302	L=13	H	6	L=11	H	7	L=5	H	2	L=15	H	11	L=15	H	12	L=12	
H	240	241	H	+	L=7	H	302	L=13	H	6	L=11	H	7	L=5	H	2	L=15	H	11	L=15</td				

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.
SIMSA. 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. & Rabinvich, 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 Carlbom, 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 $\hbar = 4.0$. 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-tetramethylisoalloxazine 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).

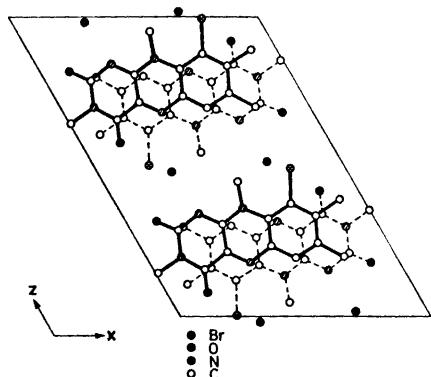


Fig. 3a. A packing diagram, excluding hydrogen atoms, viewed along the *b* axis.

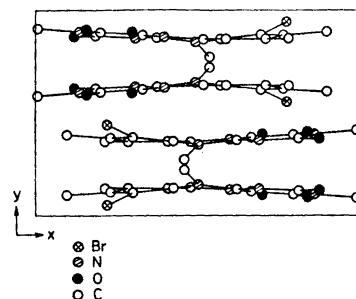


Fig. 3b. A packing diagram, excluding hydrogen atoms, viewed along the *c* axis.

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

Code for symmetry related atoms					
Superscript	Atom at	Superscript	Atom at		
None	x y z	iii			
i	$1-x$ $1-y$ $1-z$	iv	x $\frac{3}{2}-y$ $\frac{1}{2}+z$		
ii	$1-x$ $-\frac{1}{2}+y$ $\frac{1}{2}-z$	v	$1-x$ $\frac{1}{2}+y$ $\frac{1}{2}-z$		
		vi	$-1+x$ y z		
			$1-x$ $1-y$ $-z$		
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 (*c.f.* 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 resonable 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
$D = 4.2691 \text{ Å}$	C(4)	-0.078	$D = 4.1291 \text{ Å}$	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
$D = 4.0370 \text{ Å}$	C(4)	0.010	$D = 4.8325 \text{ Å}$	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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