

The Location of the Mercury Atoms in Crystals of Ribonuclease-Parachloromercuribenzoate

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The positions of the mercury atoms in ribonuclease-parachloromercuribenzoate (Ri-*pcmb*) have been determined from a study of Patterson difference projections on (010) and sharpened Patterson sections. This has permitted the calculation of the structure factors of the mercury atoms, leading to the synthesis of Fourier series for the metal-free crystal.

1. Introduction

Bernal, Carlisle and Rosemeyer (1959) showed that when crystals of ribonuclease-parachloromercuribenzoate containing chemically 1 atom of Hg per protein molecule (Dix's crystals) were soaked in 60% aqueous alcohol solutions containing 0.5% Na-*pcmb*, the symmetry of the crystals was unaffected. This diffusion process gave approximately 2 Hg atoms per protein molecule, as chemically determined by Rosemeyer. There was some evidence, from a comparison of the Patterson sections at $v = \frac{1}{2}$ for the soaked Dix's crystal and the metal-free one, that the molecules of one had moved slightly with respect to the other, showing that the crystals were not strictly isomorphous. We therefore decided to work on the soaked Dix's crystals because of the increased intensity changes in the X-ray reflections with respect to those from the metal-free one. In what follows we shall show that in the soaked Dix crystals the Hg atoms occupy two sites per protein molecule and that in the Dix crystal itself these sites are statistically filled. Since the work on the former was carried out before that on the latter, it is convenient to retain this order in the present account.

2. Location of the Hg atoms in the soaked Dix crystal

Fig. 1(a) shows a Patterson difference map on (010) (Bernal, Carlisle and Rosemeyer, 1959) using $(\Delta|F(h0l)|)^2$ between the soaked-Dix and normal crystal, incorporating reflections to spacings of 2 Å. This map was not interpretable. We next followed the procedure of Green, Ingram, and Perutz (1954) using a smoothing function of the type $\exp[-B \sin^3\theta]$, to limit the reflections to spacings of 6 Å; the resulting map is shown in Fig. 1(b). This map can be interpreted in terms of the vector relation $(A+B)^2 = A^2 + 2AB + B^2$ relating 2 pairs of atoms through a centre, which is shown graphically by the parallelogram inscribed on it. The peaks $2AB$, showing vectors between atoms not related by the centre, must fall half way between the peaks A^2 and B^2 . The additional peak seen on this

map cannot be included in this scheme and is not therefore associated with a Hg-Hg vector but is more likely to be due to a relative movement of the molecules between the two crystals.

Fig. 1(b) can only give very approximate x and z coordinates for the Hg atoms occupying sites A and B . Furthermore it is necessary to know the relative distance Δy of site A with respect to B along the b axis of the crystal (space group $P2_1$) and for this purpose we were strongly advised by Professor D. M. Hodgkin, F.R.S., to look at the individual fully-sharpened three-dimensional Pattersons of the two crystals. These calculations were very kindly carried out for us by Professor K. N. Trueblood on the Electronic Computer S.W.A.C. of the University of California.

The coefficients of the two Patterson series, each containing about 7,000 terms and scaled approximately with respect to each other, were sharpened by dividing each term by an average f^2 (averaged according to atomic percentages), multiplying by $\exp.[3B \sin^2\theta/\lambda^2]$, followed by modification with a function

$$\frac{1}{2} \sin^4\theta \cdot \exp[-50 \sin^2\theta/\lambda^2]$$

which reduces the effects of the low order terms and enhances those in the region of $3A$. The functions were calculated at 1/60 intervals of the cell edges, for the $\frac{1}{2}$ unit cell, on sections perpendicular to b , which correspond to intervals of approximately 0.5, 0.65, and 0.9 Å along a , b and c respectively. The origin peak was also removed. For economy of space and for purposes of the argument only sections $v = 6/60$, $24/60$, and $30/60$ for the two crystals have been reproduced (Figs. 2 and 3).

In the space groups $P2_1$, the equivalent points are (x, y, z) and $(\bar{x}, y + \frac{1}{2}, \bar{z})$. For 2 Hg atoms per asymmetric unit, the equivalent points are:

$$\begin{aligned} \text{Sites } A, & \quad x_1, y_1, z_1 \quad \text{and} \quad \bar{x}_1, y_1 + \frac{1}{2}, \bar{z}_1 \\ \text{Sites } B, & \quad x_2, y_2, z_2 \quad \text{and} \quad \bar{x}_2, y_2 + \frac{1}{2}, \bar{z}_2 \end{aligned}$$

Excluding the vectors at the origin, the 12 remaining vectors between these sites are of two types:

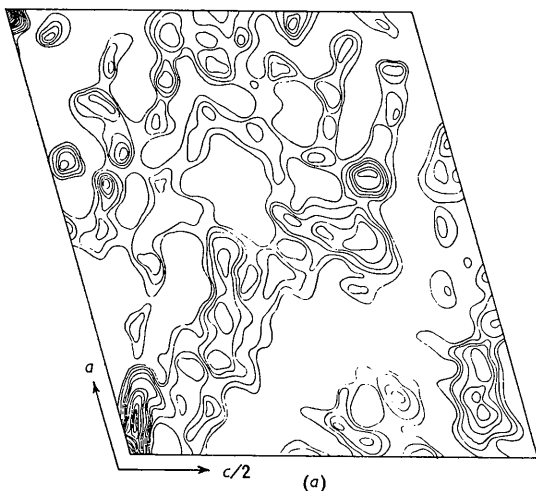


Fig. 1.

(1) Those lying on $v = \frac{1}{2}$ which may be called the A^2 and B^2 type, have components

$$2x_1, \frac{1}{2}, 2z_1 \quad \text{and} \quad \overline{2x_1}, \frac{1}{2}, \overline{2z_1}$$

$$2x_2, \frac{1}{2}, 2z_2 \quad \text{and} \quad \overline{2x_2}, \frac{1}{2}, \overline{2z_2} .$$

(2) Those of the AB type situated on the planes $v = \pm(y_1 - y_2)$ and $\pm(y_1 - y_2) - \frac{1}{2}$ having components:

$$(x_1 - x_2), \pm(y_1 - y_2), (z_1 - z_2); \quad \overline{(x_1 - x_2)}, \pm(y_1 - y_2),$$

$$\overline{(z_1 - z_2)}$$

$$\text{and } (x_1 + x_2), \pm\{(y_1 - y_2) - \frac{1}{2}\}, (z_1 + z_2)$$

$$\text{and } \overline{(x_1 + x_2)}, \pm\{(y_1 - y_2) - \frac{1}{2}\}, \overline{(z_1 + z_2)} .$$

The AB type of vectors, owing to Patterson symmetry, have mirror image relationships to each other about $v = 0$ and $v = \frac{1}{2}$; hence the appearance of vectors of the type $2AB$ when seen in projection on (010) , see Fig. 1(b).

This means that the comparisons of the Patterson sections of the crystals, with and without the metal atoms, *must be done by examining pairs of planes*, e.g. $v = 1/60$ and $29/60$, $2/60$ and $28/60$, and so on to find sufficiently heavy peaks of the AB type, of the *same height*, whose vector relationships are consistent with the appearance of the A^2 and B^2 peaks on $v = \frac{1}{2}$. It is advisable to begin the interpretation by looking for the former vectors rather than the latter because of the large number of peaks between related atoms

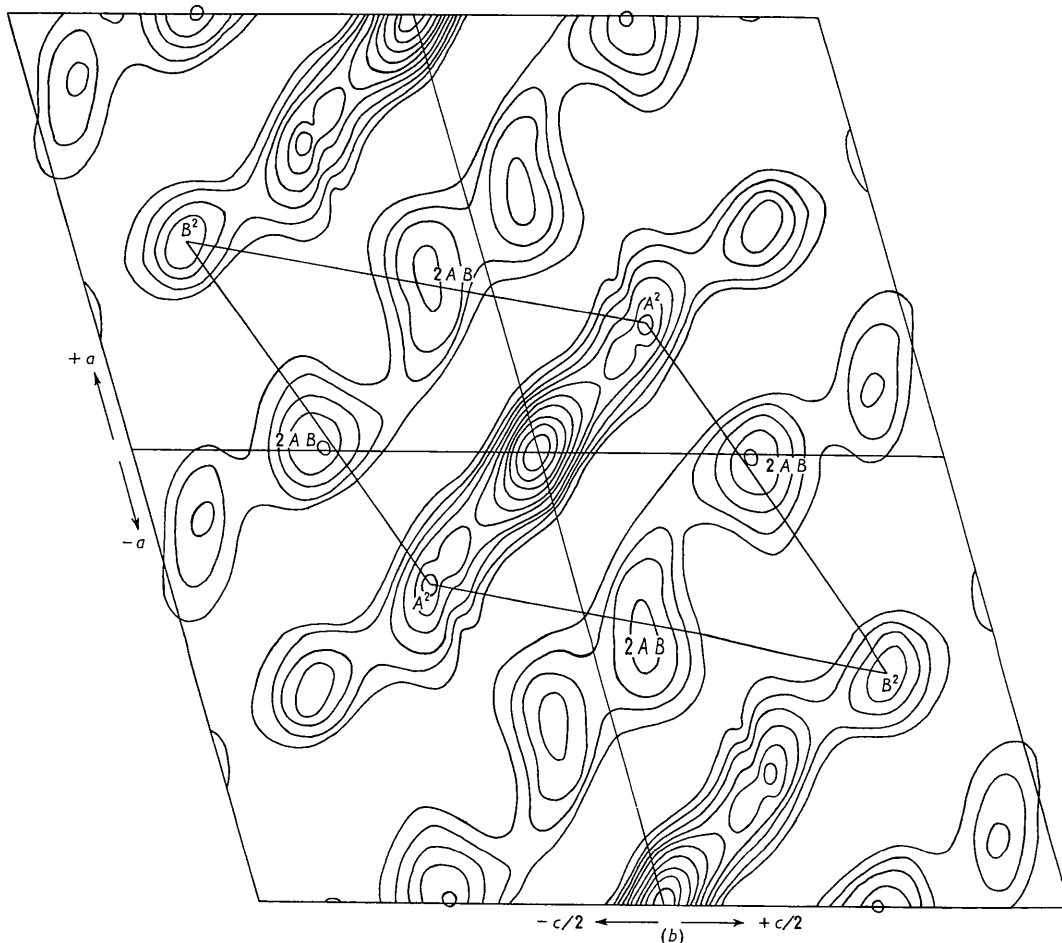


Fig. 1.

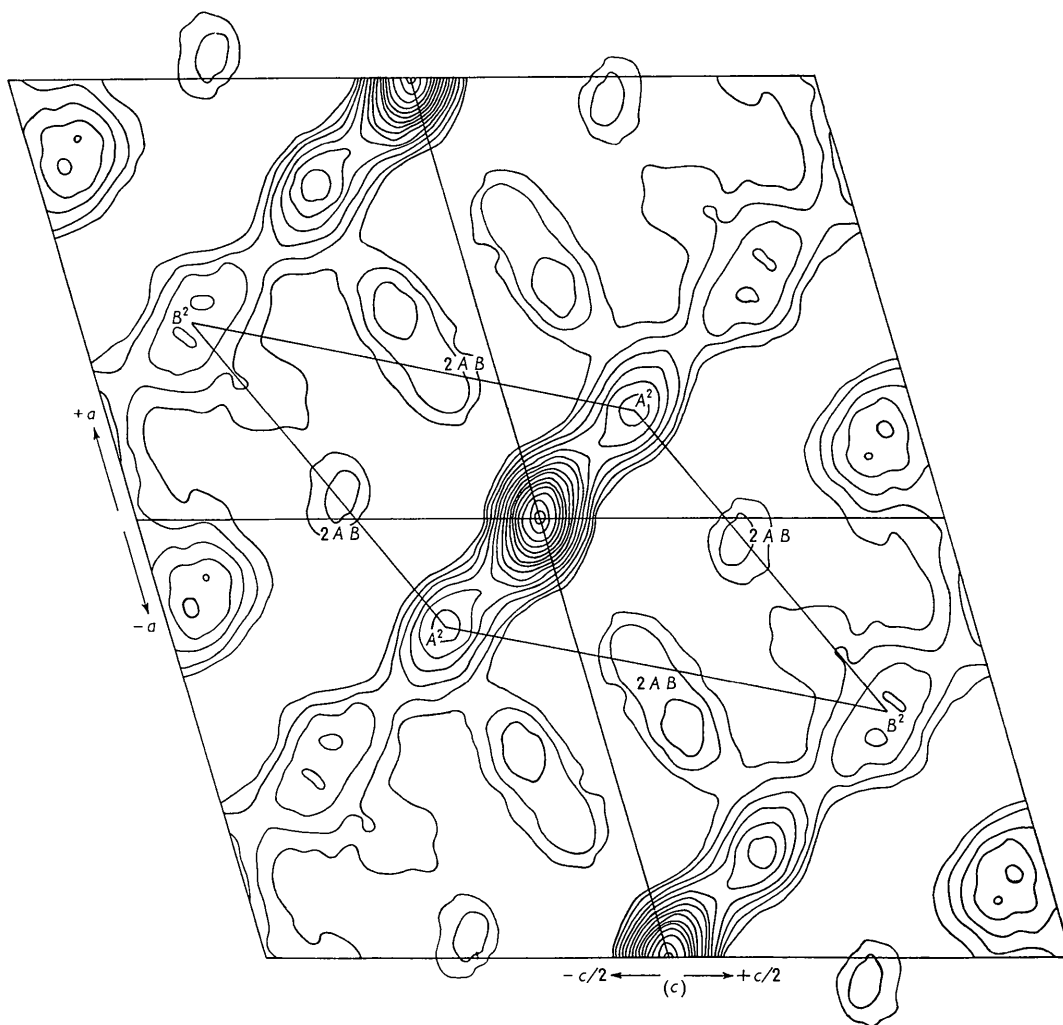


Fig. 1. Patterson difference maps projected on (010) for: (a) Soaked Dix crystal-normal crystal, using reflections with $d > 2 \text{ \AA}$. (b) Soaked Dix crystal-normal crystal, using smoothing function to terminate series at $d = 6 \text{ \AA}$. (c) Dix crystal-normal crystal, using smoothing function to terminate series at $d = 6 \text{ \AA}$.

(there being about 1000 in each molecule) which would fall on $v = \frac{1}{2}$ and so make interpretation difficult. Added to this there must also be a large number of non-Harker peaks falling on this plane, which would tend to reduce the effects of the A^2 , B^2 peaks of the Hg-Hg vectors.

If one accepts the interpretation of the Patterson difference map, Fig. 1(b), as reasonably correct, then it is only necessary to look in the $2AB$ regions of the two 3-dimensional Pattersons along b on pairs of planes equidistant from $v = 0$ and $\frac{1}{2}$ to find the AB vectors. *This was not done.* Instead, we took about half the magnitude of the average peak height for the Ri-*pcmb* Patterson function as a lower limit and any peak below this was rejected. Then we drew up a list of all peaks above this height which appeared on the Ri-*pcmb* Patterson sections but not on the corresponding Ri-sections. Fig. 4 shows a plot of these peak positions on the Patterson difference map; the number beside

each position corresponds to the section v on which it was located. For any given pair of planes, e.g., $v = 1/60, 29/60$, we took all combinations of these AB vectors, determined the x and z co-ordinates of each A and B position and attempted to locate the A^2 , B^2 vectors on $v = \frac{1}{2}$, by comparison of the corresponding Patterson maps. (Figs. 3(a) and (b)). Out of the many pairs of AB vectors that arose, there was only one that yielded vectors A^2 and B^2 that fell on positive areas on $v = \frac{1}{2}$ for the Ri-*pcmb* Patterson. In every other case either A^2 or B^2 (and sometimes both vectors) fell in negative areas on this section, thus providing sufficiently stringent conditions for their rejection.

The arrows on Figs. 2(a) and (b) indicate the AB peaks on the Ri-*pcmb* Patterson sections $v = 6/60$ and $24/60$ that were finally selected, and Figs. 2(c) and (d) show the corresponding sections for the ribonuclease crystal. The two indicated peaks on Figs. 2(a) and (b)

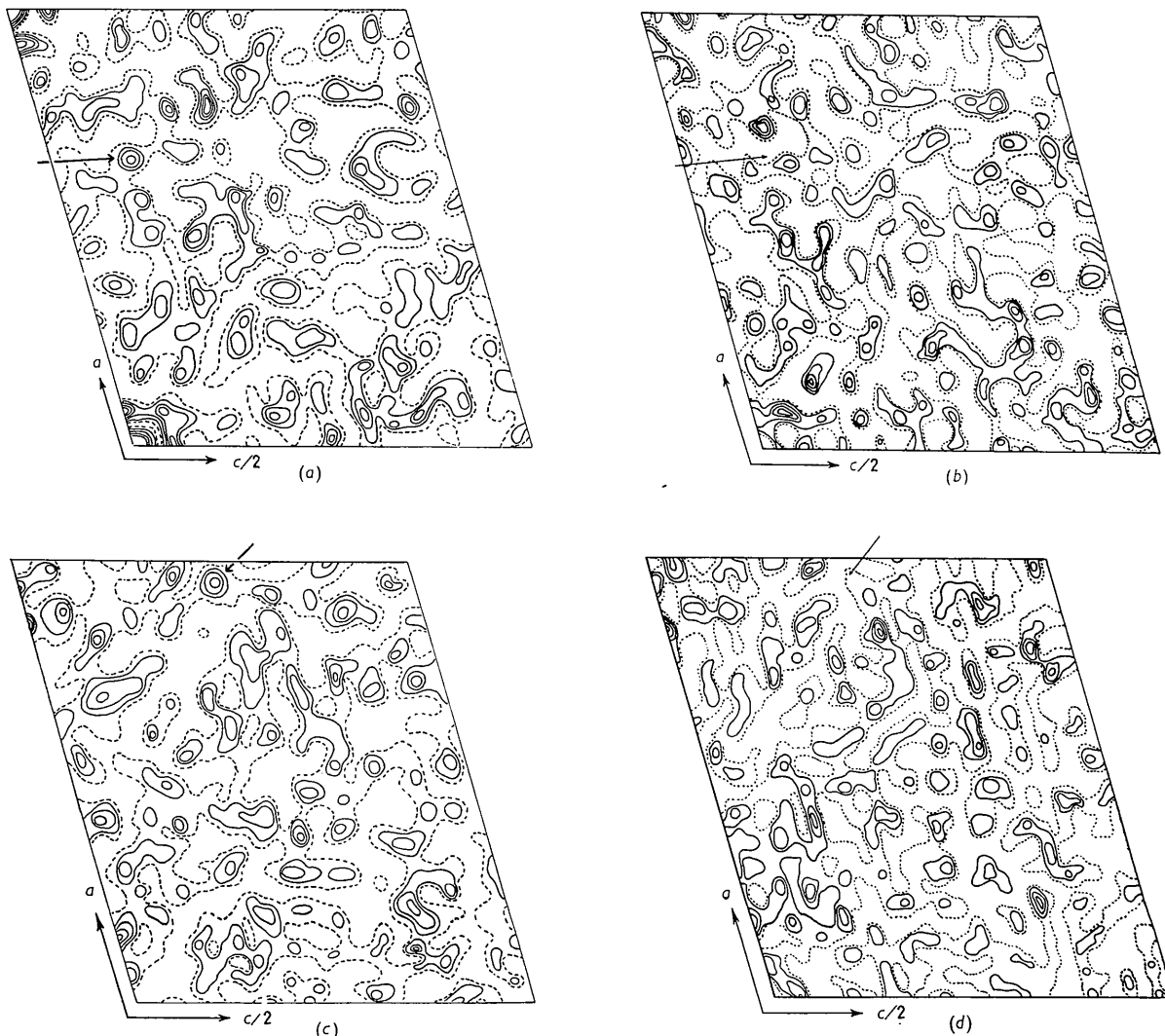


Fig. 2. Fully-sharpened Patterson sections: (a) Soaked Dix crystal, $v=6/60$. (b) Normal crystal, $v=6/60$. (c) Soaked Dix crystal, $v=24/60$. (d) Normal crystal, $v=24/60$. The arrows indicated the positions of the AB peaks of the mercury atoms in the soaked-Dix crystal.

are of the same height (and their positions have also been indicated by crosses on Fig. 4). Though they are by no means the highest in the functions, they lie well above the lower limit set in our investigation.

Figs. 3(a) and (b) show the section $v=\frac{1}{2}$ for the soaked Ri-*pcmb* and ribonuclease crystals respectively. The comparison of these two sections by themselves shows how difficult it would have been to identify the A^2 , B^2 vectors which are shown by the arrows on Fig. 3(a). The interesting and important finding is that the sharpening of the two Patterson functions has revealed the effect of the Hg-Hg vectors on this section, and for comparison we reproduce the corresponding unsharpened sections of the two crystals, Figs. 3(c) and (d). It is easily seen what sharpening has accomplished.

The AB vector on $v=6/30$ and the other on $24/30$ for the Ri-*pcmb* crystal give more reliable x and z co-ordinates for the Hg atoms than those that would have been obtained from Fig. 1(b). Using these vector relationships, the two Hg atoms in the asymmetric unit have the following co-ordinates:

	x	y	z
A	0.153	0.049	0.075
B	0.806	0.951	0.175

The y coordinates of the Hg atoms are only relative to one another and $\Delta y=0.098 \approx 6/60=1/10$. It is of interest in this connection to compare the $0k0$ reflections of the Ri- and Ri-*pcmb* crystals. This is shown in Fig. 5. The biggest changes are at 040 , $0,10,0$, and $0,12,0$. The reversal of the distribution of intensity in

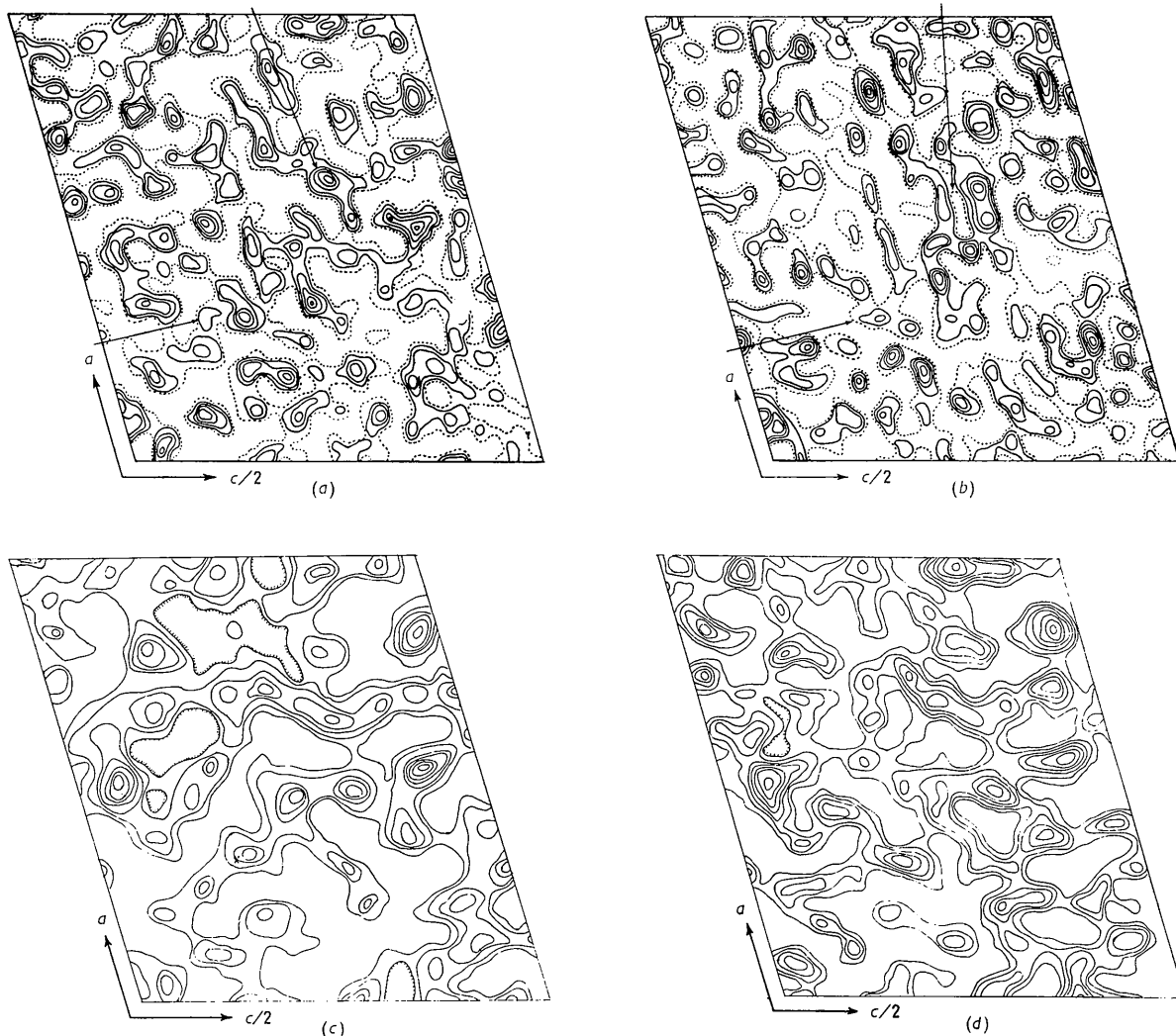


Fig. 3. Fully-sharpened Patterson sections: (a) Soaked Dix crystal, $v=30/60$. (b) Normal crystal, $v=30/60$ compared with the corresponding unsharpened Patterson sections: (c) Soaked Dix crystal, $v=30/60$. (d) Normal crystal, $v=30/60$. The arrows indicate the positions of the A^2 and B^2 peaks of the mercury atoms in the soaked Dix crystal.

going from ribonuclease to the soaked Ri-*pcmb* crystal for 0,10,0 and 0,12,0 is consistent with the Δy value found from the 3-dimensional Pattersons, and the change in 040 is due to the fact that the two Hg atoms have such a small Δy separation, and must be lying on planes approximately between the two molecules.

Table 1, column 5, shows the calculated $F(h0l)$ structure factors for the two Hg atoms to spacings of 2.5 Å. Column 4 shows the $|\Delta F(h0l)|$ of the crystals with and without the heavy atoms. As expected, the overall agreement is not very good, indicating that the relative movement of the molecules in the two crystals is possibly responsible for the discrepancies between the amplitudes of the corresponding reflections in these two columns. Some indication that this is likely to be so is shown by Fig. 6.

3. The location of the Hg atoms in the unsoaked Dix's crystal from Patterson-difference maps on (010)

Dix crystals, as stated earlier, contain 1 Hg atom per protein molecule. Since the soaked Dix crystals contain two heavy-atom sites per molecule, it is not unreasonable to suggest that the Hg atom in Dix crystal is statistically divided between the two positions. The intensities of the X-ray reflections from ribonuclease and soaked Ri-*pcmb* crystals were visually estimated. As the intensity differences of the corresponding reflections between the unsoaked Dix crystal and ribonuclease itself are smaller than the intensity differences between the soaked Dix crystal and ribonuclease for the same reflections, it is necessary

Table 1. *The calculated structure factors for the Hg atoms in the soaked Ri-pcmb crystals compared with the observed differences in structure amplitudes between the crystals with and without the metal atoms.*

<i>l</i>	$ F_{\text{Ri}} $	$ F_{\text{RiHg}} $	ΔF	$4F_{\text{Hg}}$	<i>l</i>	$ F_{\text{Ri}} $	$ F_{\text{RiHg}} $	ΔF	$4F_{\text{Hg}}$	<i>l</i>	$ F_{\text{Ri}} $	$ F_{\text{RiHg}} $	ΔF	$4F_{\text{Hg}}$
00 <i>l</i>					20 <i>l</i>					30 <i>l</i>				
1	221	491	270	214	20	0	74	74	86	0	0	104	104	-275
2	194	147	-47	0	19	127	43	-84	42	1	104	85	-19	-268
3	455	516	61	-129	18	74	134	60	-1	2	318	194	-124	-96
4	244	311	67	-94	17	199	325	126	31	3	281	345	64	78
5	147	294	147	0	16	180	0	-180	115	4	85	0	-85	112
6	85	199	114	0	15	120	321	201	147	5	120	203	83	32
7	224	134	-90	-119	14	60	0	-60	53	6	208	281	73	-19
8	180	180	0	-225	13	0	127	127	-117	7	85	180	95	49
9	216	159	-57	-181	12	336	303	-33	-221	8	85	312	227	170
10	0	0	0	0	11	357	228	-129	-174	9	0	120	120	201
11	448	285	-163	166	10	0	318	318	-38	10	265	0	-265	85
12	85	85	0	191	9	233	549	316	40	11	275	329	54	-85
13	95	112	17	93	8	398	224	174	-5	12	441	393	-48	-171
14	134	104	-30	0	7	258	443	185	-81	13	489	499	10	-125
15	127	43	-84	1	6	95	0	-95	-44	14	294	262	-32	-30
16	104	0	-104	60	5	420	258	-162	124	15	74	60	-14	9
17	224	134	-90	75	4	112	134	22	281	16	170	120	-50	-30
18	189	120	-79	0	3	332	969	637	265	17	95	85	-10	-70
19	147	95	-52	-106	2	85	60	-25	73	18	224	95	-129	-40
20	0	180	-180	-147	1	358	248	-110	-125					
10 <i>l</i>					20 <i>l</i>					40 <i>l</i>				
20	112	95	-17	-69	0	400	320	-80	-171	21	0	60	60	1
19	74	0	-74	-91	1	244	262	18	-77	20	60	208	148	46
18	175	95	-80	-20	2	104	0	-104	2	19	141	244	103	98
17	180	164	-16	111	3	60	0	-60	-50	18	120	0	-120	71
16	0	95	95	192	4	147	248	101	-172	17	159	281	122	-48
15	60	104	-44	146	5	323	405	82	-204	16	164	0	-164	-171
14	281	463	182	19	6	258	0	-258	-67	15	342	400	58	186
13	208	104	-104	-63	7	43	104	61	138	14	395	705	310	-83
12	493	794	301	-36	8	127	0	-127	240	13	164	0	-164	35
11	159	337	178	32	9	185	74	-111	173	12	337	380	43	56
10	60	159	99	15	10	417	347	-70	35	11	180	244	64	-9
9	60	368	308	-118	11	127	0	-127	-34	10	153	0	-153	-42
8	112	104	-8	-249	12	325	351	26	5	9	358	551	193	44
7	0	397	397	-226	13	221	212	-9	60	8	281	265	-16	194
6	285	258	-27	-37	14	127	199	72	30	7	164	104	-60	254
5	0	112	112	164	15	185	95	-90	-78	6	120	285	165	136
4	297	120	-177	214	16	60	74	14	-164	5	308	477	169	-73
3	0	262	262	110	17	43	85	42	-143	4	170	190	20	-196
2	441	265	-176	6	18	95	0	-95	-37	3	244	323	79	-154
1	329	236	-93	32	19	0	0	0	-58	2	224	228	4	-38
10 <i>l</i>					30 <i>l</i>					40 <i>l</i>				
0	393	515	122	145	21	134	95	-39	52	0	236	112	-124	-87
1	339	308	-31	178	20	0	43	-43	143	1	407	194	-213	-169
2	194	85	-109	35	19	170	43	-127	150	2	120	0	-120	-113
3	294	465	171	-186	18	194	216	22	58	3	203	281	78	73
4	0	303	303	-295	17	134	185	51	-51	4	104	104	0	235
5	74	208	134	-207	16	459	597	138	-79	5	85	112	27	236
6	190	95	-95	-26	15	159	0	-159	-25	6	104	43	-61	97
7	288	465	177	76	14	104	331	227	15	7	85	43	-42	-37
8	95	104	9	39	13	445	566	121	-43	8	0	95	95	-55
9	255	268	13	-35	12	294	43	-251	-162	9	0	43	43	9
10	248	424	176	-15	11	443	585	142	-207	10	268	134	-134	37
11	493	794	301	106	10	251	0	-251	-95	11	203	0	-203	-35
12	127	0	-127	205	9	112	112	0	103	12	180	0	-180	-144
13	199	0	-199	171	8	112	0	-112	225	13	281	423	142	-173
14	291	447	156	26	7	74	141	67	178	14	236	74	-163	-86
15	0	134	134	-106	6	175	0	-175	46	15	365	199	-166	43
16	0	95	95	-128	5	258	74	-184	-15	16	0	43	43	107
17	153	194	41	-60	4	719	1200	481	51	17	0	0	0	77
18	127	95	-32	-3	3	380	424	44	132					
19	185	104	-81	-16	2	221	236	15	81	50 <i>l</i>				
20 <i>l</i>					1	407	337	-70	-107	20	175	0	-175	-83
21	60	120	60	58						19	127	74	-53	-66
										18	164	85	-79	-11

Table I (cont.)

l	$ F_{Ri} $	$ F_{RiHg} $	ΔF	$4F_{Hg}$	l	$ F_{Ri} $	$ F_{RiHg} $	ΔF	$4F_{Hg}$	l	$ F_{Ri} $	$ F_{RiHg} $	ΔF	$4F_{Hg}$
										50l				
17	147	0	-147	-5	8	74	60	-14	-94	0	85	185	100	-86
16	385	358	-27	-76	9	95	251	156	-165	1	120	141	21	-107
15	0	0	0	-147	10	0	95	95	-125	2	445	399	-46	-41
14	153	300	147	-114	11	233	221	-12	4	3	268	141	-127	4
13	391	127	-264	33	12	0	112	112	114	4	275	251	-24	-40
12	244	0	-244	180	13	306	281	-25	126	5	175	43	-132	-121
11	141	473	332	203	14	0	43	43	56	6	134	185	51	-133
10	337	175	162	97	15	95	104	9	-1	7	120	60	-60	-39
9	212	120	-92	-16						8	199	159	-40	89
8	315	281	-34	-23						9	127	228	101	145
7	147	159	12	53						10	199	120	-79	99
6	203	307	104	79						11	95	95	0	17
5	0	104	104	-30						70l				
4	288	347	59	-200	20	371	486	115	-20	18	306	288	-18	-93
3	194	297	103	-269	19	0	0	0	-77	17	153	297	144	-16
2	216	104	-112	-159	18	306	288	-18	-93	16	175	268	93	111
1	153	208	55	36	17	153	297	144	-16	15	43	74	31	180
					16	127	255	128	129	14	127	255	128	129
					13	285	203	-82	14	13	285	203	-82	14
					12	194	244	50	-51	12	194	244	50	-51
					11	0	0	0	-19	11	0	0	0	-19
					10	134	0	-134	38	10	134	0	-134	38
					9	104	185	81	12	9	104	185	81	12
					8	398	291	-107	-109	8	398	291	-107	-109
					7	281	312	31	-213	7	281	312	31	-213
					6	561	809	248	-182	6	561	809	248	-182
					5	199	85	-114	-24	5	199	85	-114	-24
					4	43	0	-43	129	4	43	0	-43	129
					3	153	180	27	156	3	153	180	27	156
					2	221	281	60	72	2	221	281	60	72
					1	0	74	74	2	1	0	74	74	2
										80l				
					0	185	112	-73	33	18	74	95	21	126
					1	153	85	-68	118	17	74	74	0	45
					2	0	127	127	133	16	112	74	-38	-42
					3	0	60	60	21	15	104	60	-44	-57
					4	433	473	40	-135	14	60	60	0	-11
					5	475	342	-133	-202	13	60	0	-60	15
					6	141	439	298	-134	12	134	43	-91	-40
					7	422	0	-422	-13	11	0	85	85	-133
					8	0	153	153	46	10	120	74	-46	-160
					9	112	153	41	15	9	278	297	19	-67
					10	164	0	-164	-30	8	0	0	0	79
					11	141	134	-7	-9	7	74	0	-74	158
					12	281	85	-196	74	6	320	306	-14	117
					13	203	236	33	133	5	85	0	-85	26
										4	224	255	31	-6
										3	95	95	0	46
										2	300	236	-64	96
										1	104	141	37	53
										90l				
					0	120	95	-25	-75	0	120	95	-25	-75
					1	0	147	-147	-178	1	0	147	-147	-178
					2	104	0	-104	-164	2	104	0	-104	-164
					3	255	208	47	-54	3	255	208	47	-54
					4	0	104	104	47	4	0	104	104	47
					5	74	74	0	59	5	74	74	0	59
					6	0	0	0	10	6	0	0	0	10
					7	112	0	-112	-14	7	112	0	-112	-14
					8	104	147	43	34	8	104	147	43	34
					9	34	159	25	105	9	34	159	25	105
										100l				
					16	272	324	52	-42	16	272	324	52	-42
					15	332	199	-133	-133	15	332	199	-133	-133
					14	85	0	-85	-137	14	85	0	-85	-137
					13	0	0	0	-56	13	0	0	0	-56
					12	112	134	22	25	12	112	134	22	25
					11	127	104	-23	32	11	127	104	-23	32
					10	194	180	-14	-16	10	194	180	-14	-16
					9	147	85	-62	-33	9	147	85	-62	-33
					8	112	255	143	35	8	112	255	143	35
					7	258	342	84	138	7	258	342	84	138
					6	120	175	55	169	6	120	175	55	169
					5	255	351	96	84	5	255	351	96	84
					4	244	228	-16	-49	4	244	228	-16	-49
					3	134	74	-60	-119	3	134	74	-60	-119
					2	194	251	57	-86	2	194	251	57	-86
					1	74	43	-31	-18	1	74	43	-31	-18
										10,0l				
					0	275	255	-20	-5	0	275	255	-20	-5
					1	127	127	0	-61	1	127	127	0	-61
					2	0	104	104	-106	2	0	104	104	-106
					3	0	60	60	-65	3	0	60	60	-65

Table 1 (cont.)

<i>l</i>	$ F_{Ri} $	$ F_{RiHg} $	ΔF	$4F_{Hg}$	<i>l</i>	$ F_{Ri} $	$ F_{RiHg} $	ΔF	$4F_{Hg}$	<i>l</i>	$ F_{Ri} $	$ F_{RiHg} $	ΔF	$4F_{Hg}$
10,0 <i>l</i>					11,0 <i>l</i>					11,0 <i>l</i>				
4	216	104	-112	46	9	159	43	-116	57	0	85	60	-25	23
5	199	164	-35	135	8	258	422	164	10	1	95	0	-95	78
6	0	60	60	129	7	159	134	-25	6	2	0	95	95	52
11,0 <i>l</i>					6	0	0	0	42	3	306	228	-78	4
13	0	43	43	-79	5	203	190	-13	51	12,0 <i>l</i>				
12	112	180	68	26	4	190	112	-78	-22	8	0	43	43	-7
11	60	95	35	122	3	112	60	-52	-124	7	0	127	127	-97
10	240	95	-145	129	2	60	0	-60	-157	6	95	0	-95	-107
					1	127	134	7	-87	5	0	0	0	-48
										4	306	228	-78	1

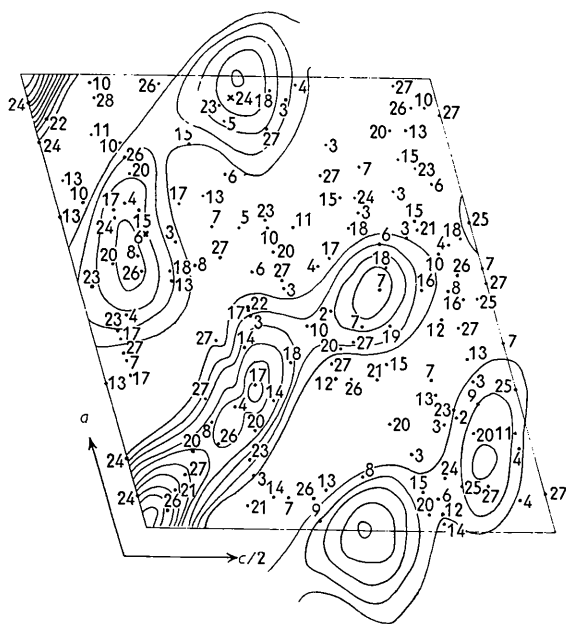


Fig. 4. Plot on Fig. 1(b) of all peaks occurring on fully-sharpened Patterson sections of soaked Dix crystal having peak height greater than half the average peak height. The crosses indicate the positions of the *AB* peaks of the mercury atoms in the soaked-Dix crystal as determined from the fully-sharpened Patterson sections.

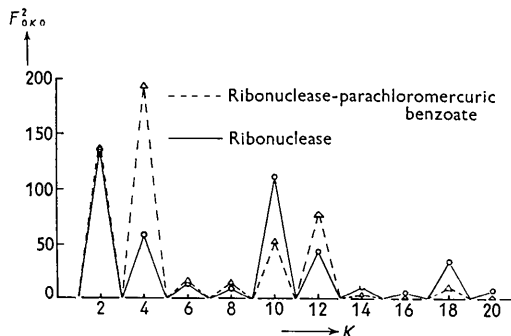


Fig. 5. Comparison of the intensities of the $0k0$ reflections between the soaked-Dix and normal crystals.

to photometer the reflections from Dix crystals and ribonuclease if a reliable conclusion is to be drawn

about Hg positions that are now making $\frac{1}{4}$ of their full contribution to the intensities of the reflections. The following work briefly describes the procedure adopted.

The investigation was confined to the $h0l$ reflections of the two crystals, whose diffraction patterns were recorded with a Buerger Precession camera which facilitated the measurement of intensities by means of a JCL-Walker Automatic Recording Microdensitometer.

Crystals of about the same size, of both types, were carefully selected for the intensity measurements, and the intensities were corrected for Lorentz-polarisation effects by Waser's function given in Volume II of the *International Tables for Crystallography*. The $F^2(h0l)$ values from the two crystals were independently scaled by the Wilson method (1942), with due allowance made for the presence of the Hg atoms in Dix's crystal. The Patterson difference map using $(\Delta|F|)^2$ as coefficients was calculated for spacings to 2.5 Å; it showed similarities to Fig. 1(a) and was also uninterpretable. A smoothing function was then used which terminated the series at 6 Å, and Fig. 1(c) shows the resulting Patterson difference map projected on (010).

The similarity between Figs. 1(b) and (c) is at once apparent, though the peaks on 1(c) show different relative heights compared with those on 1(b). The pattern however is again interpretable in terms of 2 pairs of atoms related through a centre. The x and z coordinates of the *A* and *B* sites, which can only be approximate, are given in the table below together with the corresponding x and z positions determined from the analysis of the 3-dimensional Patterson sections of the soaked crystals.

Crystals	<i>A</i> (site)		<i>B</i> (site)	
	x	z	x	z
Unsoaked Dix crystal	0.12	0.08	0.78	0.18
Soaked Dix crystal	0.153	0.075	0.806	0.150

The agreement is sufficiently close to show that the mercury atoms occupy the same sites in both the unsoaked and soaked Dix crystal. The Δy values for the *A* and *B* sites in Dix's crystal have not been

determined, but it is unlikely that they will be any different from those already found for the soaked ones.

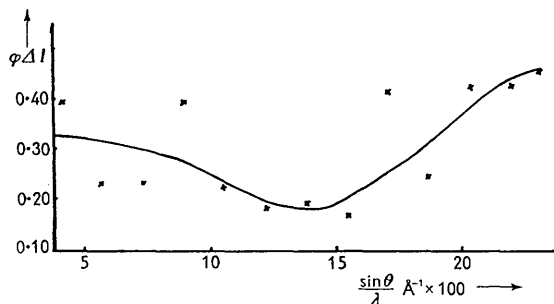


Fig. 6. Plot of the fractional change in intensity, $\phi \Delta I = 2\{(\Delta|F|)^2/\bar{I}_{R1}\}^{\frac{1}{2}}$, for Dix crystal—normal crystal, versus $\sin \theta/\lambda$. The initial fall in $\phi \Delta I$ is due to the mercury atoms in the Dix crystal and the increase for $\sin \theta/\lambda > 0.14$ is attributed to a shift in the molecules of one crystal relative to those of the other.

Fig. 6 shows a plot of $\phi \Delta I = 2\{(\Delta|F|)^2/\bar{I}_{R1}\}^{\frac{1}{2}}$ against $\sin \theta/\lambda$ (Crick and Magdoff, 1956). The initial decrease in the function is due to the effect of the heavy atoms, but the small increase towards higher $\sin \theta/\lambda$ values indicates that there must be a slight shift of the molecules in the Dix crystals relative to those in the metal-free ones. This is in agreement with the previous findings of Bernal, Carlisle and Rosemeyer (1959) for the soaked crystals.

4. Conclusions

This work has shown how the positions of heavy atoms in protein crystals, particularly when there is more than one per asymmetric unit, may be found by using fully sharpened three-dimensional Patterson functions, and this is all the more important where there is a likelihood that isomorphism is not strictly obeyed owing to the relative movement of the molecules in the two crystals.

The structure factors of the mercury atoms in the soaked crystal have now been calculated to spacings

of 2\AA , leading to the synthesis of two three-dimensional Fourier distributions for the metal-free crystal incorporating some 60% of the reflections in each case to (a) the 6\AA region, and (b) the 2.5\AA region. From this it has been possible to gain some knowledge of the configuration of the single polypeptide chain in this molecule and also the positions of the four cystine sulphurs in it.

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