tween cyclopolysilanes and aromatic hydrocarbons. Further investigations of cyclopolysilanes seem likely to provide the key to understanding of controversial questions of bonding in metalloid compounds even as studies of their carbon analogs, the cyclic and cage hydrocarbons, have been crucial to present knowledge of organic stereochemistry and reaction mechanisms.

The reactions of cyclopolysilanes are not only interesting in themselves, but have opened the way to the synthesis of complex polysilanes and thus to whole new areas of study. Improved methods of synthesis and isolation are needed, but the number and kinds of compounds that can be prepared seem almost limitless. Perhaps a polymetal chemistry comparable in breadth and variety to carbon chemistry is now developing.

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My debt goes also to my early mentors in science: Joseph Bouckaert, Joseph Maisin, Hugo Theorell, Carl and Gerty Cori, and Earl Sutherland. Four of them have preceded me on this podium. Three, unfortunately, are not with us any longer.

Exploring Cells with a Centrifuge

Christian de Duve

In one of her masterpieces, Nobel Laureate Selma Lagerlöf tells how the little boy Nils Holgersson visited the whole of Sweden, from Skåne to Lappland, on the wings of a friendly white gander. I too have made a wonderful journey, using, like Nils Holgersson, an unconventional mode of travel. For the last 25 years, I have roamed through living cells, but with the help of a centrifuge rather than of a microscope.

On these trips I was never alone. I want to mention this at the onset, since I owe much to my traveling companions. Some of their names will come up as my tale unfolds; but there are so many of them that I will be quite unable to mention them all.

The Development of Analytical **Cell Fractionation**

Thirty years ago, much of the living cell still remained virtually unexplored. The reasons for this are simple. Morphological examination was limited downward in the scale of dimensions by the resolving power of the light microscope, whereas chemical analysis stopped upward at the size of the smaller macromolecules. In between, covering almost two orders of magnitude, lay a vast terra incognita, impenetrable with the means of the day. Invasion of this territory started almost simultaneously on its two frontiers, after electron microscopy became available to morphology and centrifugal fractionation to biochemistry.

When, in 1949, I decided to join the little band of early explorers who had followed Albert Claude in his pioneering expeditions, electron microscopy was still in its infancy. But centrifugal fractionation, the technique I wanted to use, was already well codified. It had been described in detail by Claude himself (1), and had been further refined by Hogeboom, Schneider, and Palade (2), and by Schneider (3). According to the scheme developed by these workers, a tissue, generally rat or mouse liver, was first ground with a Potter-Elvehjem homogenizer, in the presence of either 0.88M (2) or 0.25M (3) sucrose. The homogenate was then fractionated quantitatively by means of three successive centrifugations and washings, under increasing centrifugal force \times time integrals, to yield "nuclei," "mitochondria," "microsomes," and a final supernatant. The fractions, as well as the original homogenate, could then be analyzed for their chemical composition, enzyme content, and other properties.

All these details were available in the literature, and there seemed little more for us to do than to acquire the necessary equipment and follow instructions carefully, especially since our interest in cell fractionation itself was rather peripheral at that time. All we wanted was to know something about the localization of the enzyme glucose 6-phosphatase, which we thought might provide a possible clue to the mechanism of action, or lack of action, of insulin on the liver cell.

Fortunately, this is not exactly how things happened. Working with me on this project was Jacques Berthet, still a medical student at that time, but with an unusually mature and rigorous mind. He went about the job of setting up the technique in a careful and systematic fashion, paying special attention to all physical parameters. A few practical tips from Claude, who had just returned to Belgium, were also helpful.

Particularly important, I now realize in retrospect, was the fact that we took some time to study the theory of centrifugation, as beautifully exposed in the classical book by Svedberg and Pedersen (4).

Although separating mitochondria and microsomes might appear worlds apart from the determination of the molecular weight of macromolecules, certain concepts were common to the two operations and could be usefully transposed-from the latter to the former. One was that of sedimentation coefficient (Fig. 1), which obviously was applicable to any particle, irrespective of its size. Another was that of .polydispersity, which, owing to biological variability, was likely to be a property of the populations made up by subcellular organelles. This meant that the centrifugal behavior of such populations could be described only by a frequency distribution curve of sedimentation coefficients (Fig. 2), not by a single s value as for most molecular populations. A third important point related to the resolving power of differential sedimentation, which some elementary calculations revealed to be surprisingly low (Fig. 3).

There was much insistence in those days on the various artifacts that complicate centrifugal fractionation, such as, for instance, breakage or agglutination of particles, and adsorption or leakage of soluble constituents. But these were only accidents, no doubt serious but amenable to experimental correction. The problem, as it appeared to us, was a more fundamental one. What we were doing was trying to separate populations that, owing to overlapping polydispersities, might at best be only partly separable from each other. In addition, we were using a poorly discriminating method for this purpose.

I cannot claim that all this was immediately clear to us. But considerations of this sort undoubtedly colored our approach from the start (5). We fully expected centrifugally isolated fractions to be impure, while suspecting that populations of cell organelles might be difficult, if not impossible, to resolve quantitatively. Conscious also of the severe limitations of light microscopic examination of the fractions, we tried to extend the biochemical interpretation as far as possible. Instead of looking at each fraction separately and focusing on its enzyme content, as was usually done, we looked rather at each individual enzyme and contemplated its distribution between all the fractions.

In order to permit a comprehensive view of enzyme distribution patterns, I introduced a histogram form of representation (Fig. 4). In this figure are shown the distribution patterns of three of the first enzymes we studied, on the left as determined by the classical four-fraction



For spherical particle of radius r (cm) and of density $ho_{
m D}$ (g × cm⁻³)

In medium of density $\rho_{\rm m}$ (g × cm^-3) and of viscosity η (poises)

s =
$$2r^2(\rho_p - \rho_m)/9\eta$$

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The author is Andrew W. Mellon Professor at Rock-efeller University, New York 10021, and professor at the Université Catholique de Louvain, Louvain, Bel-gium. This article is the lecture he delivered in Stock-holm, Sweden, on 12 December 1974 when he received the Nobel Prize for Physiology or Medicine, a prize he shared with Albert Claude and George Palade. The article is published here with the permission of the Nobel Foundation and will also be included in the complete volume of *Les Prix Nobel en 1974* as well as in the series Nobel Lectures (in English) published by the Else-vier Publishing Company, Amsterdam and New York. The lectures by Claude and Palade will appear in later

Fig. 1. The Svedberg equation and its application to a spherical particle.



Fig. 2 (left). Image of a polydisperse population of particles. Owing to individual differences in size or density (or both), different members of the population do not have the same sedimentation coefficient. The centrifugal properties of the population as a whole are depicted by a frequency distribution curve of sedimentation coefficients. Size or density distributions can be similarly represented. Frequency is usually defined as dn/Ndx (or, in the case of histograms, $\Delta n/N \Delta x$) in which (dn/N) ($\Delta n/N$) is the fraction of total particles having an abscissa value comprised between x and x + dx (Δx). Instead of relative number, similar diagrams may be drawn in terms of relative mass, relative enzyme activity, and so forth. Fig. 3 (right). The percentage of particles recovered in a sediment as a function of relative particle volume. Particle density is assumed to be the same for all particles. The meniscus of fluid in the rotating centrifuge is assumed to be halfway between the axis and the bottom of the tube or cell.





scheme, and on the right as determined by the modified five-fraction scheme that we worked out in an effort to elucidate the significance of the small difference in distribution observed between acid phosphatase and cytochrome oxidase (6). This difference, as can be seen, is very much magnified by the modification in fractionation scheme.

These histograms turned out to be very revealing, by more or less automatically conveying the notion of polydispersity, illustrated in Fig. 2. In fact, since the fractions are aligned along the abscissa in order of decreasing sedimentation coefficient, one may, in a very crude fashion, look at the abscissa as a deformed scale of sedimentation coefficients, and at the histograms as correspondingly deformed frequency distribution histograms of sedimentation coefficients. The logical next step in this line of reasoning was to assimilate enzyme distributions to particle distributions, and therefore to interpret, at least tentatively, significant differences in the distribution patterns of two enzymes as reflecting association of the enzymes with distinct particle populations.

Extrapolation from enzymes to particles could not, however, be made without some sort of assumption concerning the relationship between relative enzyme activity, the numerator in the ordinate of Fig. 4, and relative particle number, the numerator in the ordinate of Fig. 2. The simplest, and at the same time most plausible, such assumption was that members of a given particle population have essentially the same biochemical composition, larger particles simply having more of everything than smaller particles. Within the limits of validity of this assumption, which I have called the postulate of biochemical homogeneity, the histograms of Fig. 4 could now be likened to distribution diagrams of total particle mass or protein (not of actual particle numbers, it should be noted, although further conversion to numerical distributions can be made with some additional information). We had to assume, of course, that the enzyme distributions were not grossly distorted by translocation artifacts, or to correct for such artifacts as much as possible.

Another postulate we made was that each enzyme is restricted to a single intracellular site. This postulate of single location is less essential than that of biochemi-

Fig. 4. Enzyme distributions represented in histogram form. The relative specific enzyme content (percentage of activity divided by percentage of protein) of the fractions is plotted against their relative protein content, inscribed cumulatively from left to right in their order of isolation (decreasing sedimentation coefficient): nuclear (N), mitochondrial (M), microsomal (P), and supernatant (S), in classical four-fraction scheme; and nuclear (N), heavy mitochondrial (M), light mitochondrial (L), microsomal (P), and supernatant (S), in modified five-fraction scheme (6). Although very crude, the similarity with frequency distribution curves of polydisperse populations can be recognized. Distinction between three populations, now known to consist of mitochondria (cytochrome oxidase), lysosomes (acid phosphatase), and endoplasmic reticulum fragments (glucose 6-phosphatase), is enhanced by use of five-fraction scheme. [Source: (44)]

cal homogeneity, since bimodal or multimodal distributions are amenable to the same kind of interpretation. In practice, however, single location made a useful addition to biochemical homogeneity, supporting the use of enzymes as markers of their host particles.

First used empirically as pure working hypotheses, the above considerations were progressively validated, as more enzymes were studied and a limited number of typical distribution patterns began to emerge. Actually, as shown by the results of Fig. 5, things were not quite as simple, and a number of complications of various sorts tended to blur the picture. But most of these could be dealt with satisfactorily by ancillary experiments (7).

In these studies, a second line of evidence based on enzyme latency, proved very useful. Owing to impermeability of particle membranes to one or more of the substrates used in the assay of enzymes, many particle-bound enzymes fail to display activity "in vitro" as long as the membrane surrounding them is intact. Various means, mechanical, physical, or chemical, can be used to disrupt the membrane and to release the enzymes, as we first showed for rat liver acid phosphatase (Fig. 6). If two or more enzymes are present together in the same particles, they will be released together in this kind of experiment; if in different particles, they may come out separately (Fig. 7). In our hands, such studies have been very useful, providing an independent verification of the significance of the similarities and differences revealed by centrifugation experiments.

By 1955, our results were sufficiently advanced to allow us to propose with a certain measure of assurance the existence of a new group of particles with lytic properties, the lysosomes, and to hint at the existence of another group of particles, the future peroxisomes (7). At the same time, we had, from the mixture of theoretical considerations and experimental results that I have just briefly recalled, derived a certain "philosophy" of centrifugal fractionation, which I subsequently elaborated in greater detail in several publications (8). The key word here was "analytical." Basically, we felt that our approach was no more than an extension of the classical Svedberg technique from the molecular to the submicroscopic and microscopic level.

A major difficulty at this stage, however, was that available techniques did not measure up to the kind of information we were hoping to extract. The answer to this problem was provided by density gradient centrifugation, which was introduced in the early 1950's. This new technique offered prospects of improved resolution; it allowed the use of density, as well as of sedimentation coefficient, as a separation parameter; and, finally, its analytical character was unmistakable (Fig. 8). In fact, as shown as early as 1954 by Hogeboom and Kuff (9), it could even be used successfully for the determination of molecular weights.

Here again, we devoted some time to theoretical studies (10). In this, Berthet and I were joined by another young co-

worker, Henri Beaufay, whose skills as a self-taught engineer proved particularly valuable for the design of various accessories, culminating in the construction of a completely automatic rotor (11), different in principle from the zonal rotors built by Norman Anderson (12), and particularly adapted to rapid isopycnic separation at minimum hydrostatic pressure. The importance of the latter advantage has been



Total nitrogen (%)

Fig. 5. Distribution patterns of enzymes in rat liver fractions separated by five-fraction procedure shown in Fig. 4. Pattern I, shared by three enzymes, represents the distribution of mitochondria; pattern II (glucose 6-phosphatase), that of microsomes. In between, in the left column, are complex combinations of patterns I and II. Pattern III is shared by five lysosomal acid hydrolases, except for β -glucuronidase which has an additional microsomal component. Pattern IV belongs to the peroxisomal urate oxidase. The numbers of determinations are given in parentheses. Details are given in the original report. [Source: (7)]





Fig. 6 (left). Model of latency of rat liver acid phosphatase, as proposed in 1951 (58). [Source: (59)] Fig. 7 (above). Differential release of the lysosomal acid phosphatase and of the peroxisomal catalase by increasing concentrations of digitonin. [Source: (59)]

emphasized by my former collaborator Robert Wattiaux (13).

Particles sedimenting through a density gradient are apt to undergo a progressive increase in density, due to inflow of solute or outflow of water or both, depending on the number and permeability properties of their membranes and on the nature of the solute (or solutes) and solvent used to make the gradient. These factors we tried to incorporate in a theoretical model of particle behavior (10, 14), and at the same time to take into account in the design of our experiments. It appeared from our theoretical considerations that the sucrose concentration of the medium might be a particularly important variable, and that different types of particles might respond differently to changes in sucrose concentration. We therefore subfractionated large granule fractions from rat liver in isosmotic glycogen gradients prepared with sucrose solutions of different concentrations as solvent, as well as in sucrose gradients prepared with either H_2O or $D_2O(15)$.

The results of these experiments con-

firmed and extended our earlier findings, establishing the existence of three distinct groups of enzymes, as defined by their centrifugal behavior. There was little doubt in our minds that these observations reflected the occurrence of three distinct populations of particles in the large granule fraction. By fitting our results to the theoretical equation, we were even able to evaluate a number of physical parameters for each putative particle population and to construct, from purely biochemical data, a sort of "robot picture" of the particles themselves (Table 1). Due to heterogeneity within the population, the data given in this table for the lysosomes are of questionable significance. On the other hand, those listed for mitochondria and peroxisomes agree very well with measurements made by other techniques.

Although analytically satisfactory, the results described so far still fell short of definitive proof, since they had unfortunately confirmed our fear that distinct populations of subcellular particles might prove intrinsically inseparable quantitatively due

Table 1. Typical physical properties of rat liver particles [Source: (59)]

Reference enzyme Mitochondria Peroxisomes Lysosomes Parameter Cytochrome Acid Acid deoxy-Urate d-amino acid Catalase ribonuclease oxidase phosphatase oxidase oxidase Dry weight (µg) 10-7 2.7×10^{-8} 3.6×10^{-8} 2.4×10^{-8} 1.319 1.315 1.315 1 300 1.331 1 322 Drv density Osmotically active solutes 0 0 0 0.157 0.128 0.334 (milliosmole/g, dry weight) Water compartments (cm³/g, dry weight) Hydration 0.430 0.256 0.212 0.214 0 295 0.296 0.905 1.075 0.330 2.51 2.682.54 Sucrose space 0.595 0.485 1.265 0 0 Osmotic space in 0.25M sucrose 0 1.807 2.724 2.975 2.836 Total in 0.25M sucrose 1.930 1.816 Sedimentation coefficient in 0.25M 104 4.4×10^{3} 5×10^3 4.4×10^{3} sucrose (Svedberg units) 0.51 0.56 0.54 Diameter in 0.25M sucrose (µm) 0.8 1.088 1.090 1.099 1.103 1.100 1.095 Density in 0.25M sucrose

to overlapping of size or density distributions, or both. It was possible to obtain pure samples by cutting off nonoverlapping parts of the populations, but this introduced the danger of biased sampling. A means of almost complete separation, although under somewhat artificial conditions, was provided in 1962 by Wattiaux, Maurice Wibo, and Pierre Baudhuin (16), when they discovered that treatment of the animals with Triton WR-1339 causes a selective decrease in the density of subsequently isolated lysosomes, due to accumulation of the Triton within these particles (Fig. 9). Thanks to this finding and to the Beaufay rotor, large-scale separation of the three populations has now become possible, allowing a variety of biochemical and functional studies that were not feasible before (17).

While the biochemical approach I have outlined was being developed in our laboratory, electron microscopy was making great strides of its own, soon becoming available for the examination of subcellular fractions. For obvious reasons we were very anxious to take a look at our purest fractions, in order to test our conclusions and eventually identify our hypothetical particles. Already in 1955, thanks to the expert collaboration of Alex Novikoff and to the facilities of Claude in Brussels and of Wilhelm Bernhardt in Paris, we were able to do this for lysosome-rich fractions, which were found to contain dense bodies, surrounded by a membrane and of about the size predicted for lysosomes (18). Later, we were able to acquire an instrument of our own, and Beaufay taught himself another skill, which he later perfected under the guidance of George Palade. With Baudhuin, he confirmed the identification of lysosomes as "pericanalicular dense bodies" and showed that the peroxisomes correspond to the particles known as "microbodies" (19). Thus, the gap between biochemistry and morphology was finally bridged, after some 15 years of research.

More recently, Baudhuin has adapted quantitative morphometric methods to the examination of subcellular fractions, making it possible to compare measurements derived from biochemical data with those obtained by direct mensuration (20). In several instances, excellent agreement has been found between the two sets of data (17, 20-22).

Applications to Biology

I have chosen to dwell at some length on our theoretical and technical studies, because they were, I believe, the key to whatever achievements were made by our group. I know that others have accomplished important advances by the alternative process of first purifying a subcellular component and then analyzing it. For example, nuclei, secretion granules, plasma membranes, and Golgi elements have been largely characterized in this fashion. But purification is generally a laborious procedure, it is difficult to control, and it is rarely quantitative. The advantage of the analytical approach is that it is widely applicable, and it can provide a considerable amount of quantitative information, even with a relatively poor resolving power. The important point is that with this kind of methodology, we derive the information not from the properties of specific fractions believed to approximate a given intracellular component, but from the manner in which properties are distributed over a large number of fractions, which together represent the whole tissue.

In our laboratories, this general approach has been applied to a variety of biological materials and for the study of many different problems. In continuation of the work on liver, already described, it has supported a number of studies concerned with the functions of lysosomes, including those of Wattiaux on intralysosomal storage (23), of Pierre Jacques on pinocytosis (24), of Russell Deter on autophagy (25), of Jack Coffey, Nick Aronson, and Stanley Fowler on lysosomal digestion (26), and of André Trouet and Paul Tulkens on the effects of antibodies against lysosomes (27). It has also allowed Brian Poole, Federico Leighton, Tokuhiko Higashi, and Paul Lazarow to make a searching analysis of the biogenesis and turnover of peroxisomes (21, 28). In recent years, a large team grouped around Beaufay and Berthet, and including Alain Amar-Costesec, Ernest Feytmans, Mariette Robbi, Denise Thinès-Sempoux, and Wibo, has launched a major

attack on microsomal and other membrane fractions with the aim of characterizing physically, chemically, enzymically, and immunologically the various types of cytomembranes occurring in these fractions (29).

In its applications to other mammalian tissues and cell types, analytical cell fractionation has allowed Baudhuin and Poole to recognize peroxisomes in kidney (30); Gilbert Vaes to carry out a thorough study of bone lysosomes, leading to very revealing observations on the role of these particles in bone resorption (31); Bill Bowers to make a comprehensive biochemical dissection of lymphoid tissues and lymphocytes, as a preliminary to an analysis of cell-mediated immune cytotoxicity (32); Marco Baggiolini to characterize the two types of granules present in neutrophil polymorphonuclear leucocytes (33); Richard Schultz and Jacques to unravel some of the complexities of placental tissue (34); Tim Peters to fractionate aortic smooth muscle cells (35) and enterocytes (36); and Tulkens to do the same for cultured fibroblasts (37), a system also used by Poole and Wibo for investigations of protein turnover (38).

Under the leadership of Miklós Müller, a series of fascinating studies have been performed in New York on a number of different protozoa. In Tetrahymena pyriformis, Müller was able to identify two types of lysosomes, which discharge their enzymes, one in phagocytic vacuoles and the other in the outside medium (39). In collaboration with Baudhuin and later with Jim Hogg, he has shown the existence in the same organism of peroxisomes that. like plant glyoxysomes, contain enzymes of the glyoxylate cycle (30, 40). More recently, with Don Lindmark, he has characterized in Trichomonas a completely new type of cytoplasmic particle, with the capacity of converting pyruvate to acetate, carbon dioxide, and molecular hydrogen, the hydrogenosome (41).

Other studies have dealt with the role of lysosomes in tissue regression, notably those of Denise Scheib-Pfleger and Wattiaux on Müllerian ducts in chick embryos



1. Differential sedimentation

Gradient: Shallow stabilizing, pmax. < pmin.

Centrifugation: --- Incomplete sedimentation

Abscissa of frequency distribution: Sedimentation coefficient

2. Density equilibration

Gradient: Steep, p_max > p_max

Centrifugation: Prolonged, high speed

Abscissa of frequency distribution: Equilibrium density

Fig. 8. Schematic representation of density gradient centrifugation, with initial top layering of the sample. Two forms, based on differences in sedimentation coefficient and density, respectively, are shown. Diagram at the right pictures frequency distribution of particles or markers as a function of tube height. Conversion to frequency distributions of sedimentation coefficients or densities generally requires readjustment of ordinate and abscissa values, leaving surface area of each block (percentage of content in fraction) unchanged. For details of calculations, see (10). [Source: (59)]

(42), and those of Yves Eeckhout on the tail of metamorphosing tadpoles (43).

It has been my good fortune to participate in most of these investigations, sometimes actively and sometimes simply in an advisory capacity, and to watch at the same time the growing interest of other laboratories in similar problems. After trying, with increasing difficulty, to review the field of lysosomes at regular intervals (44, 45), I welcomed with some relief the appearance in 1969, under the editorship of John Dingle and Honor Fell, of the multiauthor treatise Lysosomes in Biology and Pathology (46), of which volume 4 is now in press. The literature on peroxisomes and related particles has grown more slowly, but has now also reached an appreciable size (47).

It must be pointed out that many of these advances have been made by means of morphological rather than by biochemical methods, or by a combination of both. In this respect, the development of cytochemical staining reactions for enzymes previously identified biochemically as specific particle markers has been an invaluable aid, thanks to the pioneering work of Novikoff, Stanley Holt, Werner Straus, Fritz Miller, Sidney Goldfisher, Marilyn Farquhar, and many others.

Applications to Pathology and Therapeutics

In recent years, we have become increasingly concerned with the possible medical applications of our findings. The possibility that lysosomes might accidentally become ruptured under certain conditions, and kill or injure their host cells as a result, was considered shortly after we got our first clues to the existence of these particles. We even made a number of attempts to test this hypothesis in ischemic tissue and in the livers of animals subjected to hepatotoxic treatments or to carcinogenic diets (48). But we became discouraged by problems of interpretation (45). Even today, clear-cut demonstration of the socalled "suicide bag" hypothesis remains very difficult, although there seem to be at least a few authenticated cases involving this mechanism of cell death. Much more clearly documented is the mechanism of tissue injury through extracellular release of lysosomal enzymes, a field that has been pioneered by Fell and her co-workers.

The two mechanisms mentioned above rely on the plausible instance of lysosomal enzymes exerting their lytic effect at abnormal sites. What we did not suspect in the beginning was that the failure of lysosomal enzymes to act at their normal site could also cause serious diseases. This fact was brought home to us in a rather surprising fashion through the work of my colleague Géry Hers, who in 1962 diagnosed glycogen storage disease type II as being due to a severe deficiency of a lysosomal enzyme (49). This finding initiated a



Fig. 9 (left). Influence of a previous injection of Triton WR-1339 on the equilibrium density of rat liver particles equilibrated in an aqueous sucrose gradient. Upper graph shows overlapping of lysosomes (A. Pase, acid phosphatase; A. DNase, acid deoxyribonuclease) with mito-

chondria (*Cyt. ox.*, cytochrome oxidase) and peroxisomes (*Ur. ox.*, urate oxidase). Four days after intravenous injection of 170 mg of Triton WR-1339 to the animals, the density of the lysosomes has decreased drastically, whereas that of mitochondria and peroxisomes remains unchanged; graph constructed from results of Wattiaux *et al.* (*16*). [From: (*59*)] Fig. 10 (right). Influence of cholesterol feeding on density of aortic smooth muscle cell lysosomes. Graphs show distribution patterns of enzymes after density equilibration (Fig. 8) in sucrose density gradient depicted by "staircase" on top. Starting material was a postnuclear supernatant of rabbit aortic cells, brought to a density of 1.26 and layered initially at outer edge of gradient (dotted area). Broken lines give distributions in normal preparations; solid lines, those in preparation from a rabbit showing grade IV atheroma as a result of cholesterol feeding. Note extensive shift to the left of five acid hydrolases, indicating lowered density of lysosomes due to lipid accumulation. Distribution of protein, 5'-nucleotidase (plasma membranes), and mitochondrial cytochrome oxidase (not shown) was unchanged. [Source: (*52*)]

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series of fruitful investigations on other storage diseases, in which François Van Hoof played a major part (50). It also provided useful guidelines to the chemists and pathologists who, in various parts of the world, were trying to unravel the pathogenesis of hereditary lipidoses and mucopolysaccharidoses. Today, with more than 20 distinct congenital lysosomal enzyme deficiencies identified, this mysterious chapter of pathology has been largely elucidated (51).

According to some results obtained over the last few years by Peters, Müller, Tatsuya Takano, Bill Black, and Helen Shio, with the collaboration of Farquhar, lipid accumulation in arterial cells during the development of atherosclerosis could well be due to a mechanism similar to that involved in congenital lipidoses. At least in cholesterol-fed rabbits, there is strong evidence, both biochemical and morphological, that the lysosomes of the aortic smooth muscle cells are the main site of intracellular cholesterol ester accumulation. and there are indications that a relative deficiency of the lysosomal cholesteryl esterase may be responsible for this phenomenon (35, 52, 53). Figure 10 shows some of the biochemical evidence: after cholesterol feeding, lysosomes become considerably less dense due to lipid accumulation. This figure also illustrates the sensitivity of our present techniques. These fractionations were performed on a total of about 1 milligram of cell protein. Similar experiments have been successfully performed on a needle biopsy.

Other interesting applications of the lysosome concept are in pharmacology and therapeutics. In line with the "suicide bag' hypothesis, early investigations in this area focused on "labilizers" and "stabilizers" of the lysosomal membrane (54). One outcome of this work has been the suggestion that certain anti-inflammatory agents, such as cortisone and hydrocortisone, might owe at least one part of their pharmacological properties to their effect on the lysosomal membrane.

More recently, we have extended our interest to the various substances that are taken up selectively into lysosomes and owe some of their main pharmacological properties to this phenomenon. These "lysosomotropic" agents are surprisingly numerous, including such variegated compounds as neutral red, chloroquine, streptomycin, dextran, polyvinylpyrrolidone, Triton WR-1339, and trypan blue (55). Particularly interesting is the use of certain lysosomotropic agents as carriers for drugs. In Louvain, Trouet has applied this principle to leukemia and cancer chemotherapy, by using DNA as carrier for the

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drugs daunorubicin and adriamycin. Experimentally, these DNA complexes proved less toxic and more effective on L1210 leukemia than the free drugs (56). Clinical trials under way over the last 2 years in several hospitals have given very encouraging results (57).

Conclusion

In the conclusion of his Nobel lecture delivered in 1955, Hugo Theorell asked: "What is the final goal of enzyme research?"

"The first stage," he answered, "is to investigate the entire steric constitution of all enzymes. . . . In the second stage," he continued, "it is a matter of deciding how the enzymes are arranged in the cell-structures. This implies, as a matter of fact, the filling of the yawning gulf between biochemistry and morphology.'

The gulf still yawns today. But it is a particular pleasure for me to be able to tell my old friend Theo that it yawns a little less. In our efforts to narrow it, my coworkers and I have been privileged to contemplate many marvelous aspects of the structural and functional organization of living cells. In addition, we have the deep satisfaction of seeing that our findings do not simply enrich knowledge, but may also help to conquer disease.

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will not refer to a particular intention, involved in a "whole" and in some way transcending the partial effects.

It is in this way that the "past"-that is, those aspects of past phenomena which are amenable to measurement-has come to appear as the "cause" of the present. It is a concise statement summarizing the accumulated experience obtained by observing the behavior of nonliving bodies and systems, collected since the beginning of modern science.

In this century it has become evident that in the atomic and electronic domain the measurable data are not sufficient for a completely definite prediction of succeeding states; there is dispersion in the development and statistical predictions are the most that can be made. Thus, causal relationship is partly deterministic, partly statistical. In the investigation of these relations no evidence has been found for the effectiveness of "finalistic causes," that is, causes directed to the future. All predictions that can be made are still statements which relate the present to the past. It has also become clear that in the atomic and electronic domain every observation or measurement disturbs the system under observation.

There is no need to dwell on the power the scientific method of observation has given to mankind. However, it should not be overlooked that this enormous success has also depended on the type of problems studied. Many questions occupied the human mind during the Middle Ages, for example, the problem whether man's destiny was to adjust himself to a cosmic order, embracing both the moral and the material world. The new science of Leonardo, Galileo, and their followers was directed away from such problems and substituted a new set of questions. Pushed in this new direction, Western thought came more and more to rely on the assumption that everything in the Universe is determined by what has occurred in its past. It is here that a warning note is in order. There is no justification for enforcing this concept of causality on the entire Universe as the only possible form of relationship. In particular, while many phenomena exhibited by living beings can be foreseen on the basis

Causality and Anticipation

Analysis of the concept of anticipation can contribute to the philosophy of biology.

J. M. Burgers

The purpose of this article is to renew discussion of the problem whether the phenomena of life can be satisfactorily analyzed and explained on the basis of the laws discovered in the physical sciences, or whether more is needed. When mentioning the physical sciences, I have in mind the physical laws as they are formulated at present, with the trend of thinking that forms their present background. Otherwise the problem would become indefinite. I wish to consider the thesis that the features of life involve relations not covered by the present formulation of the physical laws, relations which, although not amenable to quantitative analysis, nevertheless play a decisive part in many reactions of living organisms. The problem is, on one hand, how to put this in appropriate terms, and on the other, to analyze some consequences of the thesis. It is useful to start with a brief recapitulation of what may be called the central doctrine of the laws of physics, namely the idea of causal relationship. This will be given in the next section. The principal argument concerning the need for extension to another form of relationship is presented in the third section of the article. It is taken from features of our human mental life (1).

Causal Relationship

Our ideas concerning causal relationship are a central feature of the physical laws. It is not necessary to present an extensive description of these ideas and a summary statement will suffice.

We are convinced that all natural phenomena occurring in systems where there is no indication of life are related to the past of the systems in such a way that knowledge of any situation gives us information on which we can base more or less adequate predictions concerning subsequent situations.

"Knowledge of a situation" means the complex of data that we can obtain by making observations and measurements according to a scheme accepted and elaborated in the physical and related sciences. In making these observations we eliminate our personal involvement. We do not introduce such qualifications as "beautiful" or "ugly," "good" or "bad." Neither do we make reference to any purpose or intention; there is no reference to the future as a determining agent. In many cases we apply dissection of complicated phenomena into more simple events, in the conviction that the separate effects can be considered as self-contained. Recombination of features follows at a later stage, and although reciprocal influences are taken into account, it is assumed that these influences

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