

Quantitative Assay and Subcellular Distribution of Enzymes Acting on Dolichyl Phosphate in Rat Liver

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ABSTRACT To establish on a quantitative basis the subcellular distribution of the enzymes that glycosylate dolichyl phosphate in rat liver, preliminary kinetic studies on the transfer of mannose, glucose, and *N*-acetylglucosamine-1-phosphate from the respective ¹⁴C-labeled nucleotide sugars to exogenous dolichyl phosphate were conducted in liver microsomes. Mannosyltransferase, glucosyltransferase, and, to a lesser extent, *N*-acetylglucosamine-phosphotransferase were found to be very unstable at 37°C in the presence of Triton X-100, which was nevertheless required to disperse the membranes and the lipid acceptor in the aqueous reaction medium. The enzymes became fairly stable in the range of 10–17°C and the reactions then proceeded at a constant velocity for at least 15 min. Conditions under which the reaction products are formed in amount proportional to that of microsomes added are described. For *N*-acetylglucosaminophosphotransferase it was necessary to supplement the incubation medium with microsomal lipids. Subsequently, liver homogenates were fractionated by differential centrifugation, and the microsome fraction, which contained the bulk of the enzymes glycosylating dolichyl phosphate, was analyzed by isopycnic centrifugation in a sucrose gradient without any previous treatment, or after addition of digitonin. The centrifugation behavior of these enzymes was compared to that of a number of reference enzymes for the endoplasmic reticulum, the Golgi complex, the plasma membranes, and mitochondria. It was very similar to that of enzymes of the endoplasmic reticulum, especially glucose-6-phosphatase. Subcellular preparations enriched in Golgi complex elements, plasma membranes, outer membranes of mitochondria, or mitoplasts showed for the transferases acting on dolichyl phosphate relative activities similar to that of glucose-6-phosphatase. It is concluded that glycosylation of dolichyl phosphate into mannose, glucose, and *N*-acetylglucosamine-1-phosphate derivatives is restricted to the endoplasmic reticulum in liver cells, and that the enzymes involved are similarly active in the smooth and in the rough elements.

Biosynthesis of the core portion of *N*-glycosidically linked saccharide chains of glycoproteins involves a number of membrane-bound enzymes and glycoside derivatives of dolichyl phosphate (reviewed in reference 42). Glycoproteins of the *Asn*-glycoside type include many secretory and membrane proteins. Therefore, the subcellular distribution of the enzymes acting on dolichyl phosphate is an essential piece of information in the understanding of secretion process and membrane biogenesis.

Biochemical data reported for various subcellular preparations suggest that the enzymes involved in core glycosylation of proteins are widely distributed among the cell membranes

(10, 13, 17, 18, 19, 23–25, 35, 41). Similar specific activities have been found in rough and in smooth microsomes from rat liver for the transfer of mannose (41) and glucose (13) from nucleotide sugars to lipid in the presence of added dolichyl phosphate, for the transfer of glucose to endogenous proteins (13), and for the transfer of an oligosaccharide from a lipid precursor to endogenous proteins (35). However, mannose is transferred from labeled guanosine diphosphate (GDP)-mannose to endogenous dolichyl phosphate and protein more efficiently by rough than by smooth microsomes prepared from hen oviduct (18) or from rat liver (41). In contrast, *N*-acetylglucosamine is transferred from uridine diphosphate (UDP)-

N-acetylglucosamine to endogenous proteins more efficiently by smooth than by rough microsomes (10) (see, however, reference 13), but it is unclear whether these transferred *N*-acetylglucosamine residues are internal or peripheral. Incorporation of *N*-acetylglucosamine into lipid derivatives in the presence of exogenous dolichyl phosphate is similar (13), or fourfold lower (10), in smooth microsomes as compared to rough microsomes. It has also been claimed that enzymes of the pathway for core glycosylation of proteins occur in mitochondria (17, 19, 23–25) and in the Golgi complex (13, 19, 41) of liver cells. In addition, studies on rat spleen lymphocytes (34), hen oviduct cells (51), and mouse fibroblasts (44) suggest that proteins can be glycosylated through the pathway of lipid intermediates at the surface of intact cells.

In fact, quantitative data on the subcellular distribution of enzymes involved in the synthesis of the core portion of the saccharide chain of glycoproteins are still lacking. The meaning of sugar incorporation into endogenous acceptors is ambiguous because the values found may reflect the amount of acceptor present as well as the enzyme activity, depending on which one is limiting. Even when a lipid acceptor was added in excess, most results published so far were not truly quantitative measurements of enzyme activities, as evidenced by the nonlinear time-course of the reaction (13, 17, 23, 25, 41) or by the complex dependence of reaction rate upon the concentration of the detergent used for dispersion of the lipid reagent and membranes (9, 10, 17, 30, 35, 41). To remove these uncertainties, we have undertaken kinetic studies on the transfer of mannose, glucose, and *N*-acetylglucosamine-1-phosphate from nucleotide sugars to dolichyl phosphate in rat liver microsomes. The subcellular distribution of enzymes involved was determined by quantitative fractionation methods. In addition, various membrane preparations were examined for their ability to carry out these reactions. The results, which have been presented earlier in abstract form (46), show that mannosyltransferase, glucosyltransferase, and *N*-acetylglucosaminophosphotransferase belong to the endoplasmic reticulum membranes and are in no way restricted to the rough portions of this cell component. Neither mitochondria, nor the Golgi complex membranes that synthesize the peripheral saccharide portion of complex glycoproteins contain detectable activities of these enzymes.

MATERIALS AND METHODS

Materials

Guanosine diphosphate [U - ^{14}C]mannose (GDP- ^{14}C Man, 173 Ci/mol), uridine diphosphate [U - ^{14}C]glucose (UDP- ^{14}C Glc, 300 Ci/mol), and uridine diphosphate *N*-acetyl- ^{14}C glucosamine (UDP- ^{14}C GlcNAc, 300 Ci/mol) were purchased from the Radiochemical Centre (Amersham, U. K.). Dolichyl phosphate (grade III, 2 mg/ml solution in chloroform/methanol, 2/1, vol/vol) was obtained from Sigma Chemical Co. (St. Louis, Mo.). Tunicamycin was a generous gift of Dr. R. L. Hamill (Lilly Research Laboratories, Indianapolis, Ind.).

Tissue Fractionation

Female rats of the Wistar strain, weighing 200–250 g, were given only water for 18 h before being killed by decapitation. All procedures were done at 2–4°C. Unless otherwise stated, the sucrose solutions used were buffered at pH 7.4 with 3 mM imidazole-HCl. Liver homogenates were obtained and subjected to differential centrifugation for quantitative fractionation into nuclear fraction (N), large granules (ML), microsomes (P), and final supernate (S) under the conditions published earlier (2). Microsomes were subfractionated by isopycnic centrifugation through a linear gradient of sucrose as described by Beaufay et al. (7). In some experiments they were subjected to digitonin as described by Amar-Costesec et al. (3).

Preparation and Subfractionation of Mitochondria

The livers from three rats were processed separately. They were chopped into pieces, added with 3 ml of ice-cold solution A (70 mM sucrose, 0.21 M mannitol, 0.1 mM EDTA, 1 mM Tris-HCl, pH 7.2 [43]) per gram of tissue, and homogenized by one passage of the teflon pestle of a tissue grinder model C (A. H. Thomas Co., Philadelphia, Pa.). The homogenate was centrifuged for 10 min at 1,700 rpm in the no. 253 rotor of the DPR-6000 centrifuge (Damon/IEC Division, Needham Heights, Mass.) and refrigerated at 2°C. The sediment was washed by two cycles of resuspension in solution A and centrifugation. Mitochondria were obtained by centrifuging the pooled supernates for 5 min (acceleration time included) at 12,500 rpm in the no. 30 rotor (Beckman Instruments Inc., Spinco Division, Palo Alto, Calif.). After decantation, the pellets were washed twice by resuspension in solution A and centrifugation; they were finally transferred into a Dounce homogenizer (Kontes Glass Co., Vineland, N. J.), added with a solution (3 ml/g of liver) of 20 mM sodium phosphate, pH 7.2, and 0.02% (wt/vol) bovine serum albumin (43), and suspended by means of the loose-fitting pestle. After 30 min at 0°C, the swollen mitochondria were given six strokes of the tight-fitting pestle to dissociate the outer membranes from the mitoplasts, and centrifuged for 20 min at 20,000 rpm in the Beckman no. 30 rotor. After decantation, the sediment was washed by resuspension in 0.25 M sucrose and centrifugation, and finally resuspended in 0.25 M sucrose. This preparation, called disintegrated mitochondria, was subfractionated by centrifugation for 60 min at 39,000 rpm in the E-40 rotor (4), loaded with 10 ml sample corresponding to ~10 g of liver, 32 ml of a sucrose gradient extending linearly with respect to the volume from density 1.10 to 1.27, and 6 ml sucrose solution of density 1.32. Subfractions were recovered and processed for density measurement and analysis as described earlier (7).

Golgi Complex Preparation

Golgi elements were prepared by following the procedure described by Wibo et al. (58).

Preparation of Plasma Membranes

Plasma membranes were prepared according to Song et al. (49), as described in a previous paper (58). Samples were withdrawn for analysis at various stages of the purification procedure. In short, the preparations consisted in membranes spun down at 1,800 rpm for 20 min in the IEC rotor no. 259 (low speed sediment), brought to equilibrium in a sucrose gradient at a density <1.18 U, and washed and resuspended in 0.25 M sucrose (type I preparation). These membranes were further purified taking advantage of the density perturbation of plasma membranes caused by digitonin (22). 1 vol of 0.25 M sucrose containing 0.75 mg digitonin/ml (~0.3 mg digitonin/mg protein) was added dropwise at 0°C under continuous stirring. A 15-ml aliquot of the suspension was laid over a step gradient loaded in the SW 25.2 Beckman rotor, and made of the following layers of sucrose solutions: 8 ml at 52.2%, 15 ml at 41.2%, 15 ml at 39.3% and 5 ml at 35.5% (wt/wt), corresponding to 1.25, 1.19, 1.18, and 1.16 density U, respectively. After 10 h of centrifugation at 24,000 rpm the membranes floating at the 1.19–1.25 interface were collected to give the type II plasma membrane preparation.

Biochemical Determinations

Aliquots of subcellular fractions and membrane preparations were quickly frozen at –80°C and thawed immediately before the assays.

Mannosyltransferase, Glucosyltransferase, and *N*-acetylglucosamine-phosphotransferase Assays

The dolichyl phosphate solution was diluted 20 times in chloroform/methanol (2/1, vol/vol), and 200- μ l portions were stored at –20°C in teflon-stoppered culture tubes (Kimax; Kimble Div., Owens-Illinois, Inc. Toledo, Ohio). Immediately before use, 0.5 μ mol EDTA and 1 μ mol $MgCl_2$ (mannosyltransferase and glucosyltransferase) or $MnCl_2$ (*N*-acetylglucosaminophosphotransferase) were added in each tube (8). The content of the tubes was thoroughly mixed and evaporated at ambient temperature under a stream of nitrogen. The dried deposit was resuspended in 50 μ l of Triton X-100 (20 mg/ml), and added with 150- μ l assay medium before starting the reaction with 50- μ l enzyme (~100 μ g of protein). The 250- μ l incubation medium contained 80 μ g/ml dolichyl phosphate, 4 mg/ml Triton X-100, 2 mM EDTA, 2.5 mM dithiothreitol, 1 mM ATP to minimize enzymic hydrolysis of the nucleotide sugars (10, 54), and the additional reagents given below.

Mannosyltransferase was assayed at 10°C and pH 7.5 in the presence of 80 mM Tris-HCl buffer, 5.8 μ M GDP- ^{14}C Man and 11.5 mM $MgCl_2$. For glucosyl-

transferase, the assay conditions were 17°C and pH 6.5, with 80 mM 2-(*N*-morpholino)ethane sulfonic acid (MES)-KOH buffer, 3.3 μM UDP-[¹⁴C]Glc and 11.5 mM MgCl₂. *N*-acetylglucosaminophosphotransferase was assayed at 17°C and pH 7.5 in the presence of 80 mM MES-glycylglycine-KOH, 28 μM UDP-[¹⁴C]GlcNAc, 10 mM MgCl₂, 4 mM MnCl₂, and heat-denatured (5 min, 100°C) microsomes from 3 mg liver. Radioactivity was ~500,000 cpm/assay. After 10 min, the reaction was stopped by adding 2.5 ml chloroform/methanol (3/2, vol/vol) and 0.4 ml of 4 mM MgCl₂. After mixing thoroughly, the phases were separated by centrifugation; the aqueous upper layer was removed, and the lower layer plus the insoluble material was washed twice with 1.25 ml chloroform/methanol/4 mM MgCl₂ (1/16/16, vol/vol/vol). An aliquot of the lower phase was assayed for radioactivity.

Chromatographic Analysis

Another aliquot of the lower phase was chromatographed on silica gel plates (Kieselgel 60, Merck A.G., Darmstadt, W. Germany) with chloroform/methanol/water (65/25/4, vol/vol/vol) as developing solvent (48). Radiolabeled glycolipids produced from the endogenous dolichyl phosphate of liver rough microsomes after incubation with UDP-[¹⁴C]GlcNAc or GDP-[¹⁴C]Man in the presence of GTP (28, 29) were used as standards.

To determine the amount of the nucleotide precursor left after incubation, the aqueous phase was analyzed by descending chromatography on Whatman no. 1 paper in ethanol/1 M acetic acid adjusted at pH 3.8 with ammonia (2/1, vol/vol).

Other Biochemical Determinations

Protein (40), NADH- and NADPH-cytochrome *c* reductases, cytochrome *c* oxidase, glucose-6-phosphatase, alkaline phosphodiesterase I, and galactosyltransferase (6), monoamine oxidase (59), *N*-acetylglucosaminyltransferase (with ovalbumin acting as acceptor [58]) and RNA (21) were assayed according to the methods described in the quoted articles. Assay of 5'-nucleotidase was performed in a 4-ml medium containing 50 mM Tris-HCl buffer, pH 7.4, 2 mM Na-AMP, 8 mM MgCl₂, and 25 μg/ml Triton X-100. After 20-min incubation at 37°C, the reaction was stopped by addition of 0.8 ml 30% TCA, the protein precipitate was removed by centrifugation, and inorganic phosphate was determined on 2-ml aliquot (20).

RESULTS

Enzyme Kinetic Studies

Kinetic studies were conducted to find out optimum assay conditions for *N*-acetylglucosaminophosphotransferase, mannosyltransferase, and glucosyltransferase, and to determine the range within which reaction velocity is proportional to enzyme concentration. Microsomes were used as a source of enzyme. The results obtained may be summarized as follows.

(a) Triton X-100 augmented noticeably the enzyme activities. However, at 37°C, its concentration in the incubation medium of mannosyltransferase was extremely critical in agreement with results from others (41, 57). A sharp peak of activity was observed at 1 mg/ml (Fig. 1*a*). The inhibition noted at higher concentrations was due to denaturation of the enzyme protein. This effect is strongly temperature-dependent, as shown by the time-course of mannosose transfer at 37°C, 25°C, and 13°C (Fig. 1*b*). In the presence of 2 mg Triton X-100/ml, the enzyme was completely inactivated after 4 min at 37°C; it was somewhat more stable at 25°C and, consequently, more mannosose could be transferred despite the lower initial velocity of the reaction at this temperature; at 13°C the reaction proceeded almost linearly for 16 min. In view of these results, mannosyltransferase was subsequently assayed at 10°C. Under these conditions, the activity increases with the concentration of Triton X-100 up to a plateau reached at 2–3 mg detergent/ml (Fig. 1*c*), and corresponding to 40-fold the activity measured in the absence of detergent, an enhancement which is much higher than that found at 37°C (Fig. 1*a*, see references 41, 57). Glucosyltransferase and *N*-acetylglucosaminophosphotransferase also lose activity at 25°C or above in the presence of Triton X-100

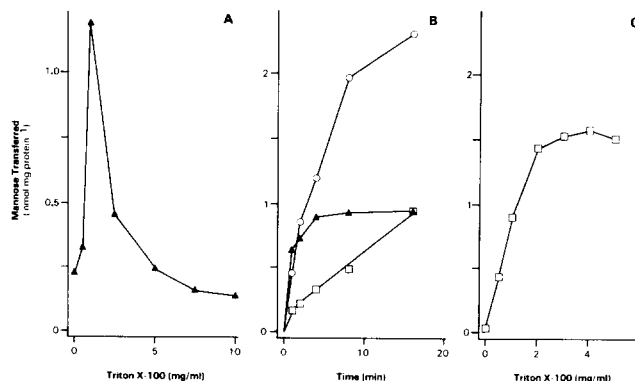


FIGURE 1 Dependence of mannosyltransferase activity on the concentration of Triton X-100 and time-course of the reaction at various temperatures. A and B, preliminary experiments in which the 250-μl incubation medium contained Tris-HCl, pH 7.4 (40 mM), GDP-[¹⁴C]Man (5.8 μM, 500,000 cpm), dolichyl phosphate (80 μg/ml), EDTA (2 mM), MgCl₂ (11.5 mM), MnCl₂ (2.5 mM), KCl (30 mM), dithiothreitol (2.5 mM), ATP (1 mM), phosphoenolpyruvate (10 mM), pyruvate kinase (2.5 U), and the microsomes from 10 mg of liver. Reactions were run at 37°C for 15 min in the presence of Triton X-100 at the concentrations given in abscissa (A), or at 13°C (□), 25°C (○), and 37°C (▲) for various times and in the presence of 2 mg/ml Triton X-100 (B). C, microsomes from 2.5 mg of liver were incubated under the conditions given in Materials and Methods (10 min at 10°C) with Triton X-100 at the concentrations given in abscissa.

(results not shown). These reactions proceed linearly for >15 min at 17°C, the temperature at which the enzymes were currently assayed. Their dependence on the Triton X-100 concentration was similar to that shown for mannosyltransferase in Fig. 1*c*, leading to 20- and 40-fold activations for *N*-acetylglucosaminophosphotransferase and glucosyltransferase, respectively.

(b) As shown in Fig. 2, the transfer reactions were influenced by the concentration of added dolichyl phosphate in a complex manner. Glucosyltransferase and, to a lesser extent, *N*-acetylglucosaminophosphotransferase (see also reference 32) activities were slightly inhibited at high dolichyl phosphate concentration. At the concentration used in the standard assay (80 μg/ml), the amount of label transferred was 40-fold (*N*-acetylglucosamine-1-phosphate), 60-fold (glucose), and 500-fold (mannose) greater than in the absence of added lipid. Such enhancement factors are far above the values currently reported (8, 17, 26, 33, 52, 57).

(c) The dependence of reaction rates upon the nucleotide sugar concentration is shown in Fig. 3. *K_m* values of 1.0 and 0.8 μM were calculated for GDP-Man and UDP-Glc, respectively. The non-Michaelian relationship observed for UDP-GlcNAc suggested that two enzymes of different *K_m* might be involved. Potentially, UDP-GlcNAc could react with dolichyl phosphate according to the following sequence:

(i) Dolichyl phosphate + UDP-GlcNAc → dolichyl pyrophosphoryl *N*-acetylglucosamine + UMP.

(ii) Dolichyl pyrophosphoryl *N*-acetylglucosamine + UDP-GlcNAc → dolichyl pyrophosphoryl *N,N'*-diacetylchitobiose + UDP.

However, the reaction did not proceed beyond formation of the monoglycoside derivative (see below), even at the highest concentration of UDP-GlcNAc used in this study. The kinetics shown in Fig. 3*b* is thus characteristic of the phosphotransferase reaction. In most fractionation experiments reported below, the activity was assayed at low (0.5 μM) and high (28 μM)

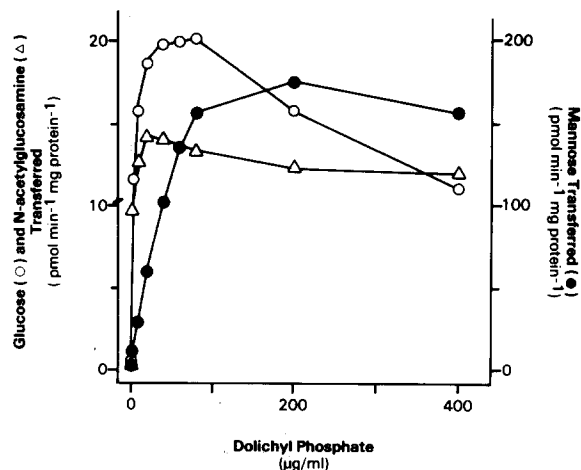


FIGURE 2 Dependence of transferase activities on the concentration of dolichyl phosphate. Mannosyltransferase (●), glucosyltransferase (○), and *N*-acetylglucosaminophosphotransferase (△) activities of microsomes derived from 2.5 mg liver were assayed at various concentrations of dolichyl phosphate. Other conditions were as given under Materials and Methods.

concentration of UDP-GlcNAc, but this did not evidence any difference in the distribution patterns (not shown). In addition, tunicamycin abolished the activity at either concentration of the nucleotide sugar, although more tunicamycin was required in the presence of 28 μ M UDP-GlcNAc to achieve the same inhibition (not shown). Consequently, if distinct *N*-acetylglucosaminophosphotransferases occur in the liver cells, they have identical subcellular distributions, and they are similarly sensitive to the antibiotic.

(d) Influence of pH was studied using Na cacodylate-HCl (pH 5.0–7.5), glycylglycine-KOH (pH 7.5–9.5), glycylglycine-MES-KOH (pH 7.0–9.5), Tris-HCl (pH 7.0–9.0), and MES-KOH (pH 5.5–7.0) buffers. Glucosyltransferase and *N*-acetylglucosaminophosphotransferase activities differed noticeably according to the buffer used (not shown); optimal conditions are those given in Materials and Methods.

(e) Owing probably to the presence of EDTA in the incubation medium, and in agreement with previous reports (8, 16, 26, 37, 55), the enzyme activities were undetectable in the absence of added divalent cations. In each case $MgCl_2$ was a more efficient activator than $MnCl_2$. The two salts have been included in the incubation medium for *N*-acetylglucosaminophosphotransferase because its activity has been demonstrated in the presence of Mg^{++} (36, 56) or Mn^{++} (10, 26, 31, 53).

(f) The reaction of mannosyltransferase and glucosyltransferase proceeded at a rate proportional to the amount of microsomes added up to ~ 8 and 4 mg liver (~ 0.4 and 0.2 mg microsomal protein), respectively (Fig. 4a and c). The relationship was also linear up to 4 mg liver when heat-denatured microsomes were included in the incubation medium of *N*-acetylglucosaminophosphotransferase reaction (Fig. 4b); otherwise, it consistently diverged from linearity in the range of 0–2 mg liver, where the activity was unexpectedly low. Addition of total lipid extracted from liver microsomes restored the expected level of activity, whereas the addition of bovine sera albumin had no effect (not shown). The added lipid could contain endogenous dolichyl pyrophosphoryl *N*-acetylglucosamine that acts as substrate for the next glucosyltransferase. We rule out this possibility, because tunicamycin completely

inhibited the reaction, and owing to the behavior of the labeled products in thin layer chromatography (see below). Most likely, some microsomal phospholipids are required for full activity of the enzyme. A curve, concave upwards, was obtained previously with solubilized enzyme preparations from hen oviduct (36) and *Acanthamoeba castellanii* (56), and with rat lung membranes (45). In the latter case addition of acidic phospholipids linearized the plot of activity vs. enzyme concentration.

For identification of the reaction products, the chloroform/methanol extracts were analyzed by thin layer chromatography. Radioautography revealed a single component, which ran with dolichyl phosphoryl mannose after incubation with GDP-

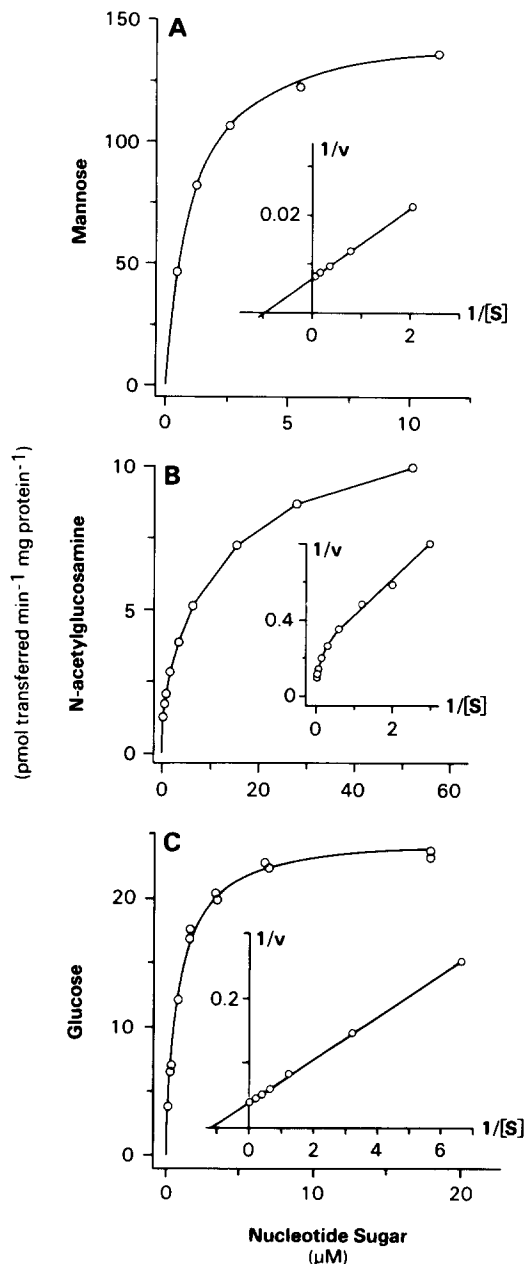


FIGURE 3 Dependence of transferase activities on nucleotide sugar concentration. Mannosyltransferase (A), *N*-acetylglucosaminophosphotransferase (B), and glucosyltransferase (C) activities of microsomes derived from 2.5 (A and C), or 0.5 mg liver (B) were assayed at various nucleotide sugar concentrations. Other conditions were as described under Materials and Methods.

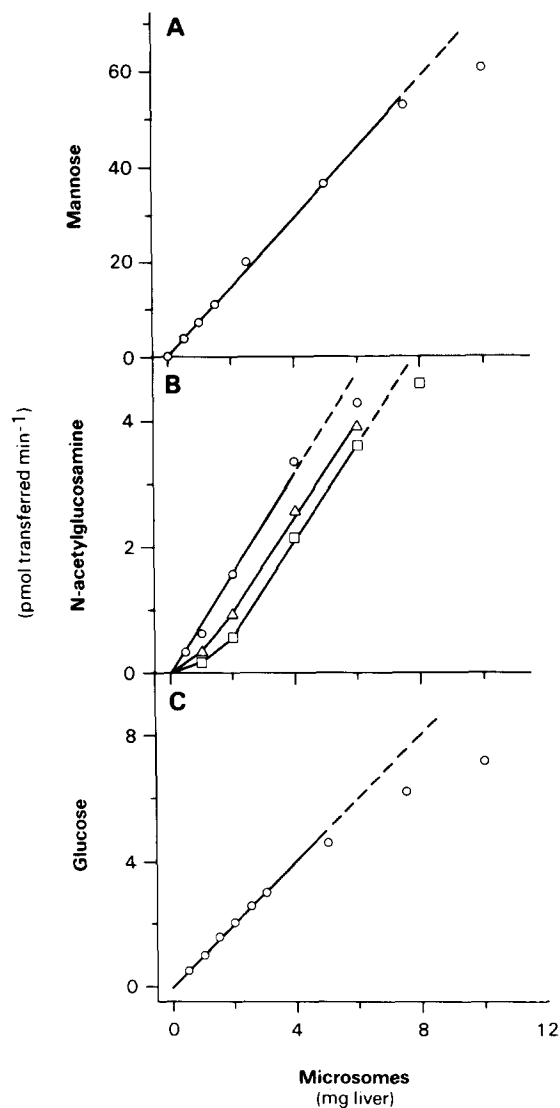


FIGURE 4 Dependence of transferase activities on amount of microsomes. Mannosyltransferase (A), *N*-acetylglucosaminophosphotransferase (B), and glucosyltransferase (C) were assayed at various microsome concentrations, as described under Materials and Methods. Reaction mixtures for *N*-acetylglucosaminophosphotransferase contained 0 (\square), 1 (Δ), or 3 mg (O) (liver weight equivalent) of heat-denatured microsomes (5 min at 100°C).

$[^{14}\text{C}]\text{Man}$, or $\text{UDP}-[^{14}\text{C}]\text{Glc}$. Mannose and glucose derivatives of dolichyl phosphate have the same mobility in this chromatographic system (33, 57). The bulk of labeled products formed upon incubation with $\text{UDP}-[^{14}\text{C}]\text{GlcNAc}$ had the chromatographic properties of dolichyl pyrophosphoryl *N*-acetylglucosamine. The amount of dolichyl pyrophosphoryl *N,N'*-diacetylchitobiose represented <5% of the label extracted, allowing to take the total label extracted as a satisfactory measure for the *N*-acetylglucosaminophosphotransferase reaction. However, the fraction of label present as a diglycoside derivative became greater upon incubation for >10 min, or at 25–37°C. Aliquots of the medium were also analyzed by paper chromatography to establish the extent of hydrolytic cleavage of the nucleotide sugars during incubation. Except for the soluble fraction, it did not exceed 10% and had no effect on the measurement of transferase activities.

Quantitative Distribution after Subcellular Fractionation by Differential and Isopycnic Centrifugation

Enzyme activities in liver homogenates and their distribution among fractions obtained by differential centrifugation are given in Table I and Fig. 5. The distribution patterns of galactosyltransferase, 5'-nucleotidase, and glucose-6-phosphatase, taken as markers for the Golgi complex, plasma membranes, and endoplasmic reticulum, respectively, are similar to those previously published (2). Other markers, including mitochondrial enzymes, were also assayed (results not shown) and behaved as in this earlier study. *N*-acetylglucosaminophosphotransferase, mannosyltransferase, and glucosyltransferase show nearly identical distribution patterns, being recovered in microsomes to the extent of ~75% of total activity. Their behavior closely resembles that of glucose-6-phosphatase.

The microsomal character of an enzyme being no proof of its association with endoplasmic reticulum, microsomes have been subjected to isopycnic centrifugation in a linear gradient of sucrose. The transferases acting on dolichyl phosphate were very similar in their density distribution (Fig. 6), but clearly differed from galactosyltransferase and 5'-nucleotidase. In agreement with earlier results (7), glucose-6-phosphatase and NADPH cytochrome *c* reductase, which both belong to the rough and smooth endoplasmic reticulum, are not identical in their density distribution, reflecting the biochemical heterogeneity of this cell component. The transferases acting on dolichyl phosphate fit better the density profile of glucose-6-phosphatase than that of NADPH cytochrome *c* reductase. This is particularly true for *N*-acetylglucosaminophosphotransferase.

Because the density distributions of the various markers overlap in the low density portion of the gradient (1.10–1.20), the data of Fig. 6 do not rule out the association of part of the enzymes acting on dolichyl phosphate with microsomes derived from the cell surface, or the Golgi complex. To examine this problem, microsomes were exposed to digitonin under conditions which do not disaggregate the membranes, and were brought subsequently to equilibrium in a sucrose gradient. Digitonin treatment causes the cholesterol-rich membranes (plasma membranes, and, to a lesser extent, Golgi complex membranes) to shift towards higher densities, presumably by digitonin binding to accessible cholesterol molecules, whereas microsomal elements derived from endoplasmic reticulum are not shifted (3). Comparison of enzyme distributions in digitonin-treated (Fig. 7, solid lines), as opposed to control microsomes (Fig. 7, shading), illustrates the shift of plasma membranes and Golgi complex elements, represented by alkaline phosphodiesterase I and galactosyltransferase, respectively. In contrast, the transferases which glycosylate dolichyl phosphate belong to digitonin-insensitive microsomal membranes.

Activity in Various Subcellular Preparations

To seek for the possible occurrence of the transferases acting on dolichyl phosphate in membranes other than those of the endoplasmic reticulum, various membrane preparations were isolated and assayed for these enzyme activities. The morphological characteristics of similar membrane preparations have been published elsewhere (58).

Golgi complex preparations appeared purified, judging from

TABLE I
Quantitative Distribution of Enzymes Acting on Dolichyl Phosphate after Differential Centrifugation

Constituent	Total activity in liver <i>nmol min⁻¹/g tissue</i>	Activity in subcellular fractions				Recovery
		N	ML	P	S	
		% of activity in total liver				
Mannosyltransferase	8.62 ± 0.99	10.1 ± 1.6	7.9 ± 3.1	78.3 ± 3.3	4.1 ± 1.9	100.4 ± 6.3
Glucosyltransferase	1.11 ± 0.17	11.6 ± 1.6	9.0 ± 3.2	80.7 ± 5.2	4.1 ± 2.1	105.4 ± 6.7
N-Acetylglucosaminophosphotransferase	0.79 ± 0.13 <i>mg/g tissue</i>	12.3 ± 0.8	11.0 ± 3.8	76.5 ± 2.8	3.6 ± 0.8	103.4 ± 4.7
Protein	245.5 ± 25	16.4 ± 0.8	24.4 ± 2.6	20.1 ± 1.4	36.8 ± 3.3	97.7 ± 3.5

Statistics refer to the mean ± SD in six experiments (five for N-acetylglucosaminophosphotransferase).

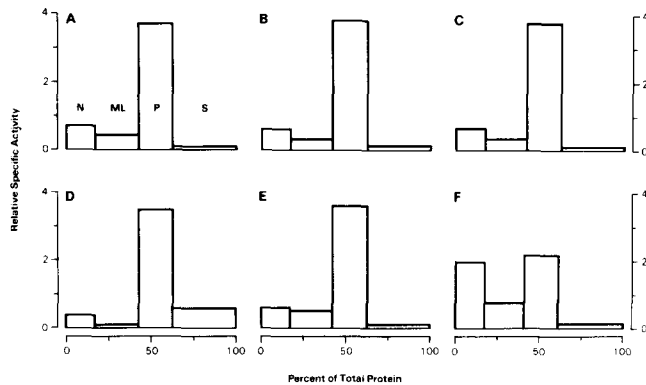


FIGURE 5 Distribution of the transferases acting on dolichyl phosphate, compared to that of reference enzymes, after fractionation of liver by differential centrifugation. These graphs were constructed as described in reference 4 from the average results of three to six experiments. Complementary data are given in Table I. Fractions are plotted from left to right in the following order: nuclear fraction (N), large granules (ML), microsomes (P), and final supernate (S). Each fraction is represented separately in the ordinate scale by the relative specific activity of the enzyme, i.e., the percentage amount of total recovered activity divided by the percentage amount of total recovered protein. In the abscissa scale, the protein content of fractions is represented cumulatively. N-Acetylglucosaminophosphotransferase (A), mannosyltransferase (B), and glucosyltransferase (C) were assayed on samples of particulate fractions which contained 40–50 μ g phospholipid. The other enzymes shown for comparison are galactosyltransferase (D), glucose-6-phosphatase (E) and 5'-nucleotidase (F), which gave average recoveries of 79%, 101%, and 101%, respectively.

the 80-fold increase in specific activity for galactosyltransferase and N-acetylglucosaminyltransferase acting on ovalbumin, when compared to total liver (Table II). For the transfer of mannose, glucose, and N-acetylglucosamine-1-phosphate to dolichyl phosphate, the relative specific activities were much lower (0.8–1.7), and comparable to those found for NADPH cytochrome *c* reductase and glucose-6-phosphatase (0.7–1.6).

Table II also shows the properties of plasma membrane preparations at different stages of their isolation: the low speed sediment, the type I preparation, and the final preparation after treatment by digitonin and isolation of the material that has acquired a higher density (type II fraction). The relative specific activities of 5'-nucleotidase and alkaline phosphodiesterase I increase in this order; those of the other enzymes, including the transferases acting on dolichyl phosphate, decrease, the final levels being 1–5% of the relative specific activity of alkaline phosphodiesterase I. The transferases acting on dolichyl phosphate are in a twofold excess over glucose-6-

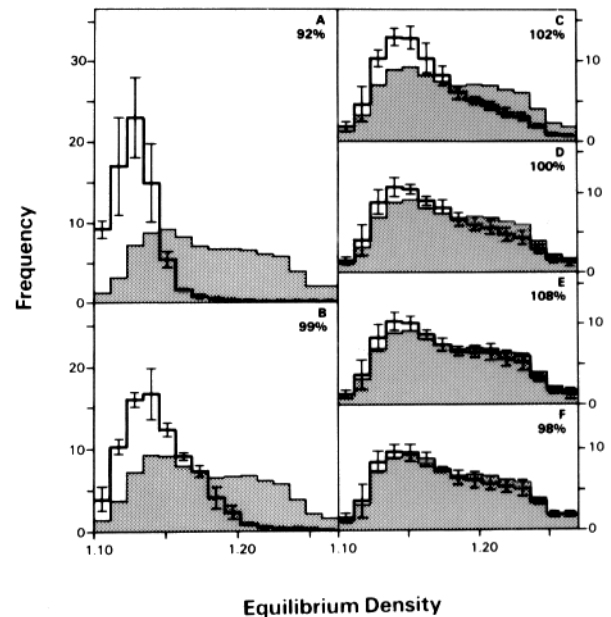


FIGURE 6 Density distribution of microsomal transferases acting on dolichyl phosphate and comparison with reference enzymes. Average results of three or four experiments in which microsomes have been centrifuged to equilibrium in a linear gradient of sucrose, as described under Materials and Methods. Frequency histograms have been normalized and averaged (see reference 4). The represented portion of histograms, divided into 15 normalized fractions of identical density increment, extends from 1.10 to 1.27 and includes >95% of the enzyme activities. Vertical lines through histogram bars give standard deviations. Solid lines are galactosyltransferase (A), 5'-nucleotidase (B), NADPH cytochrome *c* reductase (C), glucosyltransferase (D), mannosyltransferase (E), and N-acetylglucosaminophosphotransferase (F). The distribution of glucose-6-phosphatase (shading) is superimposed on each plot to facilitate comparison. Percentage values give the average recoveries from the microsome fraction. Recovery of glucose-6-phosphatase was 101%.

phosphatase and NADPH cytochrome *c* reductase at all stages.

Other experiments were devised for settlement of the question as to whether the enzymes of interest occur in the outer or inner mitochondrial membranes (Table III and Fig. 8). Mitochondria prepared by differential centrifugation were successively subjected to osmotic swelling, gentle mechanical disruption, and density equilibration in a linear gradient of sucrose. In these experiments the transferases acting on dolichyl phosphate behaved like glucose-6-phosphatase, with respect to their yield in the mitochondrial preparations, which ranged from 6.5 to 8.5% as compared to 64–69% for monoamine oxidase and cytochrome *c* oxidase (Table III), and to their density distri-

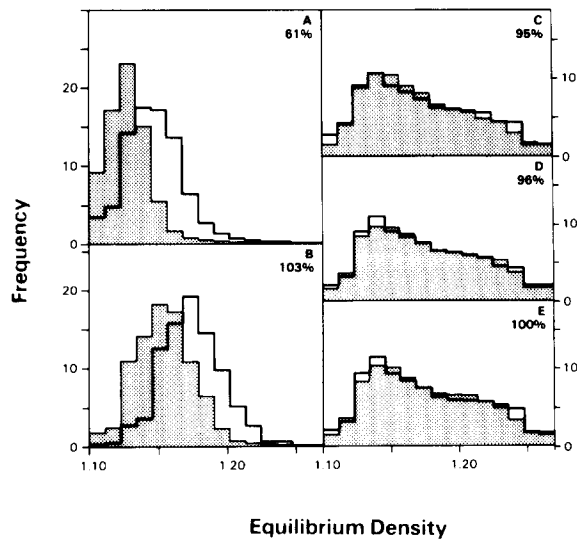


FIGURE 7 Effect of digitonin on the density distribution of the microsomal transferases acting on dolichyl phosphate. Microsomes have been treated with digitonin according to reference 3 before density gradient analysis which was carried out exactly as described in Fig. 6. Enzymes are galactosyltransferase (A), alkaline phosphodiesterase I (B), glucosyltransferase (C), N-acetylglucosaminophosphotransferase (D), and mannosyltransferase (E). The distribution obtained after digitonin treatment (solid lines) is compared to that of the same enzyme in untreated microsomes (shading, redrawn from Fig. 6, except for alkaline phosphodiesterase I). Enzyme activities of the microsome fraction were not significantly modified by digitonin treatment. Percentage values give the recovery of enzyme activities in the gradient fractions, which was 104% for glucose-6-phosphatase.

TABLE II
Enzyme Activities in Plasma Membrane and Golgi Complex Preparations

Constituent	Plasma membranes			Golgi complex
	Low speed sediment	Type I	Type II	
		% of the total liver content		
Protein	9.45	0.99	0.45	0.12
	Relative specific activity			
Mannosyltransferase	1.23	1.11	1.03	0.88
Glucosyltransferase	1.29	1.15	0.83	0.76
N-Acetylglucosaminephosphotransferase	1.32	1.21	1.08	1.67
Galactosyltransferase	ND	ND	ND	84.0
N-Acetylglucosaminyltransferase	ND	ND	ND	77.0
Alkaline phosphodiesterase I	5.04	16.6	22.2	1.4
5'-Nucleotidase	4.11	15.4	21.0	ND
NADPH cytochrome c reductase	0.71	0.85	0.48	0.74
Glucose-6-phosphatase	0.86	0.65	0.61	0.157
Cytochrome c oxidase	0.97	1.17	0.31	0.03

Subcellular fractions were obtained as described under Materials and Methods. Relative specific activities are as defined in the legend of Fig. 5. The percentage amount of enzyme activity recovered in a subcellular preparation is given by the product: percent yield of protein times relative specific activity. ND, not determined.

TABLE III
Enzyme Activities in Mitochondrial and Submitochondrial Preparations

Constituent	Disintegrated mitochondria	Subfractionation of density 1.12	Subfractionation of density 1.23
	% of the total liver content		
Protein	17.3	0.56	6.9
	Relative specific activity		
Mannosyltransferase	0.38	1.03	0.09
Glucosyltransferase	0.41	1.31	0.08
N-Acetylglucosaminophosphotransferase	0.48	1.21	0.11
		0.64	
Cytochrome c oxidase	4.0		4.4
Monoamine oxidase	3.7	57.0	0.26
NADH cytochrome c reductase	0.90	8.1	0.15
Glucose-6-phosphatase	0.49	1.22	0.11

Average results of two experiments. Relative specific activities are as defined in the legend of Fig. 5; see also Table II for estimation of the yield of enzyme activities.

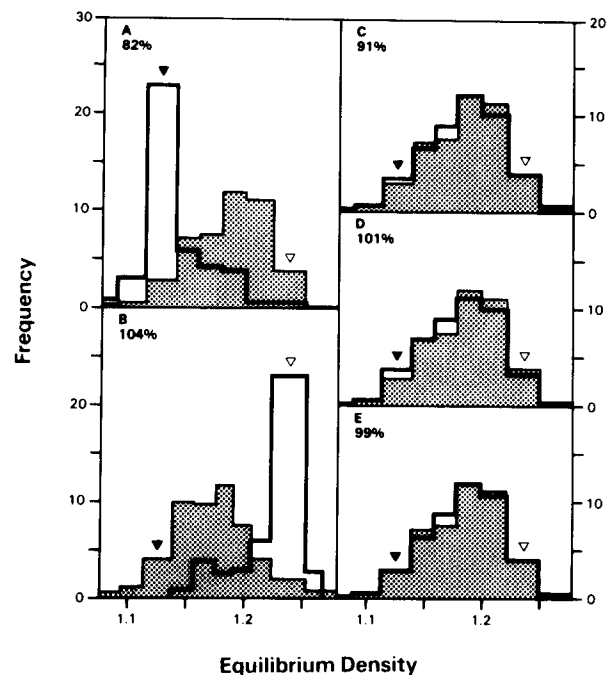


FIGURE 8 Density distribution of the transferases acting on dolichyl phosphate, compared to that of several reference enzymes, in a mitochondrial fraction subjected to hypotonic conditions and equilibrated in a sucrose gradient. Conditions for isolation of mitochondria, disintegration in a hypotonic medium, and isopycnic centrifugation are given under Materials and Methods. The represented portion of frequency histograms extends from 1.08 to 1.28 and corresponds to >95% of the enzyme activities in the mitochondrial preparation. Solid lines are monoamine oxidase (A), cytochrome c oxidase in a distinct experiment (B), mannosyltransferase (C), glucosyltransferase (D), and N-acetylglucosaminophosphotransferase (E). The distribution of glucose-6-phosphatase (shading) is superimposed on each plot to facilitate comparison. Quantitative biochemical data on the mitochondrial fraction used, and on the properties of the gradient subfractions characteristically enriched in outer (▼) and inner (▽) mitochondrial membranes are reported in Table III. Percentage values give the recovery of enzyme activities in the gradient fractions. The recovery of glucose-6-phosphatase was 98%.

bution (Fig. 8). These distributions are broad and centered around the middle of the gradient; they clearly differ from those of monoamine oxidase and cytochrome *c* oxidase, which sharply peak at modal densities of 1.12 and 1.23 U, respectively. Quantitative data on the corresponding subfractions are reported in Table III. It is apparent from these data that light (density 1.12) and heavy (density 1.23) subfractions are fairly purified outer and inner membranes of mitochondria, respectively. Like glucose-6-phosphatase, the transferases acting on dolichyl phosphate reach relative specific activities that are only 2–2.5% of those of monoamine oxidase in the outer membrane-rich fraction, and of cytochrome *c* oxidase in the inner membrane-rich fraction. In contrast, NADH cytochrome *c* reductase, an established constituent of both the endoplasmic reticulum and the outer membrane of mitochondria (50) reveals a distinctly higher specific activity in the light subfraction.

DISCUSSION

The enzymes forming the monoglycoside derivatives of dolichyl phosphate are highly susceptible to denaturation in the presence of a detergent. This property, exemplified for mannosyltransferase by the bell-shaped curve of the activity at 37°C when the concentration of Triton X-100 is varied (Fig. 1*a*), and by the nonlinear time-course at 25°C or above (Fig. 1*b*), is also apparent from other studies (17, 35, 41). We have found that mannosyltransferase, glucosyltransferase, and *N*-acetylglucosaminophosphotransferase activities can nevertheless be accurately measured provided the temperature be kept low enough, i.e., ~10–17°C. In the case of *N*-acetylglucosaminophosphotransferase, the presence of enough phospholipid in the reaction medium is an additional requirement. Our assay methods differ from those used in other laboratories only in these few aspects which turned out to be essential. The fair recovery values after subcellular fractionation (see Table I and Figs. 6–8) strengthen the kinetic evidence (Fig. 4), and argue for the validity of the methods used and of the distributions obtained.

According to these distributions, the three enzymes may be assigned to subcellular elements derived from the endoplasmic reticulum. They closely follow markers of this cell component in differential centrifugation (Fig. 5), and when microsomes or mitochondrial preparations are subfractionated by density equilibration in a linear gradient of sucrose (Figs. 6 and 8). In addition, their host membranes in the microsome fraction are unaltered in their equilibrium density after treatment with digitonin (Fig. 7). This differentiates these microsomal membranes from the Golgi complex and plasma membrane elements.

In agreement with our analytical data, several subcellular preparations enriched in either the Golgi complex, the large plasma membrane sheaths derived from the surface of hepatocytes (see 58), or the outer or inner membranes of mitochondria contain only a very small part of the total activity of the enzymes acting on dolichyl phosphate. We have recently discussed the significance of the low activities of endoplasmic reticulum-associated enzymes that are found in various preparations enriched in other subcellular components (58). The conclusion is that it reflects cross-contamination by membranes biochemically undistinguishable from those derived from the endoplasmic reticulum. This conclusion holds for the transferases acting on dolichyl phosphate because their yields are similar to those of glucose-6-phosphatase, or NADPH cyto-

chrome *c* reductase (Tables II and III). However, the three transferases are in slight excess over the reference enzymes of the endoplasmic reticulum in our plasma membrane preparations (Table II), which might be a clue in favor of their occurrence at the cell surface. Nevertheless, this possibility will not be considered, because upon increasing purification of plasma membranes the specific activity of the enzymes acting on dolichyl phosphate decreases in parallel to that of the authentic enzymes of endoplasmic reticulum.

It is thus most likely that the enzymes which form the monoglycoside derivatives of dolichyl phosphate are restricted to the endoplasmic reticulum in liver cells. This conclusion conflicts with studies reporting activity of mannosyltransferase (13, 17, 23, 41), glucosyltransferase (13, 17, 19, 24), or *N*-acetylglucosaminophosphotransferase (10, 13, 17, 25) in various other subcellular membrane entities, including the mitochondria and Golgi apparatus. The main reason for this discrepancy is that the quantitative comparison with other reference enzymes, from which our conclusion is drawn, was not possible in these earlier works. For the mitochondrial membranes, our results illustrate how misleading the presence of enzyme activities in a single membrane fraction may be (Table III). The enzyme distributions in the density gradient (Fig. 8), from which the outer and inner membrane fractions were obtained, make it evident that some enzyme activities in these fractions reflect contamination by a particular subcellular component, broadly distributed through the gradient, and peaking halfway between the two kinds of mitochondrial membranes. Contamination of mitochondrial or submitochondrial fractions by endoplasmic reticulum elements probably remained unnoticed, or underestimated in other works (17, 19, 23, 24, 25). For the Golgi apparatus, the reason for the discrepancy is perhaps also in the structural heterogeneity of this membrane complex and its close relationship with the endoplasmic reticulum. Our statement is that the enzymes acting on dolichyl phosphate, like other enzymes of the endoplasmic reticulum, occur in membranes distinct from those which carry the terminal glycosyltransferases. As discussed in detail elsewhere (5, 58), the real limits of these membrane domains are still uncertain, and may not coincide with the endoplasmic reticulum–Golgi complex junction as it is currently established on a morphological basis.

The confinement of the transferases acting on dolichyl phosphate to the endoplasmic reticulum implies that proteins glycosylated via the dolichol pathway must go through, or depend on this cell component at some stage of their biosynthesis. This is easily envisioned for glycoproteins that are synthesized by membrane-bound ribosomes. Several studies carried out on intact cells (11, 27, 38), or on cell-free systems that synthesize proteins in the presence of microsomal membranes (12, 14, 39, 47), have shown that sugar moieties are attached to various nascent chains, demonstrating that core glycosylation of proteins may occur as a cotranslational event at the level of the rough endoplasmic reticulum. According to our results, the transferases acting on dolichyl phosphate are present in both the rough and smooth portions of the endoplasmic reticulum. They markedly differ from RNA in their density distribution (see reference 7) and compare better with glucose-6-phosphatase than with NADPH cytochrome *c* reductase. The biochemical heterogeneity within the endoplasmic reticulum reflects the fact that some enzymes, e.g., glucose-6-phosphatase, assume nearly random distribution through the whole endoplasmic reticulum, whereas others, e.g., NADPH cytochrome *c* reduc-

tase, are distinctly more prominent in the smooth than in the rough portions (discussed in reference 1). Thus, the transferases acting on dolichyl phosphate belong to the first category, or even occur in slight excess in the smooth endoplasmic reticulum. This is in keeping with the finding that smooth microsomes from rat liver effectively glycosylate the nascent α subunit of human chorionic gonadotropin during synthesis in an heterologous cell-free system (15). Besides, some proteins appear to be glycosylated after release of the completed peptide chain from polysomes (12). Together with the information obtained from these other works, our results keep open the possibility that some proteins be glycosylated while they are in transit in the rough or in the smooth portions of this subcellular compartment. Quantitative studies on the other enzymes of the dolichol pathway for protein glycosylation, particularly on the enzyme that transfer the oligosaccharide from the lipid donor to the protein, may provide some clue to this matter.

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