



## Sialyl Le<sup>x</sup> Structures In *O*-Glycans Attached To Lysosomal Membrane Glycoproteins, Lamp-1 And Lamp-2. Comparison to Those in *N*-Glycans.

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**Abstract:** Poly-*N*-acetylglucosamine extension has been found in *O*-glycans in addition to *N*-glycans and glycosphingolipids (Fukuda, M. (1994) In *Molecular Glycobiology*, (ed. M. Fukuda and O. Hindsgaul), pp. 1-52, Oxford University Press, Oxford). However, the extent of poly-*N*-acetylglucosamine formation between *O*-glycans and *N*-glycans in a given glycoprotein has not been determined. Attempts were made in HL-60 cells to determine the amount of sialyl Le<sup>x</sup> structures in *O*-glycans and *N*-glycans attached to the major lysosomal membrane glycoproteins, lamp-1 and lamp-2. Lamp molecules were immunoprecipitated from <sup>3</sup>H-glucosamine labeled HL-60 cells. Glycopeptides were prepared by pronase or trypsin digestion, and *O*-glycan containing glycopeptides were isolated by affinity chromatography using Jacalin-Agarose. The glycopeptides bound to Jacalin-Agarose were treated with alkaline borohydride, and the released *O*-glycans were fractionated by Bio-Gel P-4 filtration. Similarly, the glycopeptides unbound to Jacalin-Agarose, which represent *N*-glycans, were isolated by Sephadex G-50 gel filtration. Sequential glycosidase digestion of the *O*- and *N*-glycans, with or without pretreatment by fucosidase or neuraminidase, revealed the following features: (1) each lamp-1 and lamp-2 molecule contains about 2 and 1 poly-*N*-acetylglucosaminyl *O*-glycan chains, (2) about 65% of these poly-*N*-acetylglucosaminyl *O*-glycans contain sialyl Le<sup>x</sup> termini, NeuNAcα2→3Galβ1→4(Fucα1→3)GlcNAcβ1→R, and (3) *N*-glycans and *O*-glycans are modified in almost equal efficiency to express sialyl Le<sup>x</sup> structures. These results indicate that in lamp molecules both *O*-glycans and *N*-glycans contribute to the expression of sialyl Le<sup>x</sup> structures.

Polyglucosaminoglycans are high molecular weight carbohydrates bound to proteins and lipids. Polyglucosaminoglycans are unique in having *N*-acetylglucosaminyl repeats (Galβ1→4GlcNAcβ1→3)<sub>n</sub> and this repeating structure can be identified by its susceptibility to endo-β-galactosidase. Polyglucosaminoglycans present in *N*-glycans and glycolipids have been shown to carry various antigenic determinants such as *I*<sub>i</sub> and ABO blood group antigens (for review, see Refs. 1-3). The structures of polyglucosaminoglycans can be characteristic to different cell types and stages of differentiation. In particular, it has been demonstrated that human granulocytes and monocytes express Le<sup>x</sup> structure, Galβ1→4(Fucα1→3)GlcNAcβ1→R and sialyl Le<sup>x</sup> structure, NeuNAcα2→3Galβ1→4(Fucα1→3)GlcNAcβ1→R at the termini of polyglucosaminoglycans.<sup>4,5</sup> These unique structures were found to serve as ligands for adhesive molecules, selectins, in endothelial cells and platelets,<sup>6-9</sup> and interaction between selectins and cell surface carbohydrates of granulocytes and monocytes are thought to play critical roles in the recruitment of leukocytes to inflammatory sites.<sup>10,11</sup>

Although poly-*N*-acetylglucosamines present in *N*-glycans and glycolipids have been extensively studied, it is known that *O*-glycans can also be modified by poly-*N*-acetylglucosamine extension.<sup>12,13</sup> We have shown previously that some of poly-*N*-acetylglucosaminyl *O*-glycans isolated from granulocytes can contain sialyl Le<sup>x</sup> structures.<sup>12</sup> It was also found that poly-*N*-acetylglucosamine is almost exclusively extended from the branch attached to C-6 of *N*-acetylglucosamine.<sup>12,13</sup> However, it is not known as yet which proteins have *O*-glycans containing poly-*N*-acetylglucosamines in these cells, nor how many sialyl Le<sup>x</sup> structures are present in *O*-glycans compared to those on *N*-glycans.

We have isolated two distinct groups of major sialylglycoproteins in leukocytes. Leukosialin contains approximately 80 *O*-glycans and one *N*-glycan per molecule.<sup>14</sup> It has been shown that the carbohydrate structures of these *O*-glycans are characteristic of cell lineage and different stages of differentiation.<sup>15,16</sup> We have shown that leukosialin contained, on average, one molecule of sialyl Le<sup>x</sup> terminal structure when it was isolated from HL-60 cells.<sup>17</sup> The second group consists of lysosomal membrane glycoproteins, lamp-1 and lamp-2, which contain 18 and 16 *N*-glycans, respectively, and a significant portion of these carry poly-*N*-acetylglucosamines.<sup>18,19</sup> Moreover, we found that lamp-1 and lamp-2 contain 5 and 8 *O*-glycans per molecule, respectively.<sup>20</sup>

These results prompted us to investigate the structures of *O*-glycans attached to lamp molecules, focusing on the structures of poly-*N*-acetylglucosaminyl *O*-glycans and the presence of sialyl Le<sup>x</sup> terminal structures.

## EXPERIMENTAL PROCEDURES

**Metabolic Labeling of HL-60 and K562 Cells** - For metabolic labeling of HL-60 cells, the cells were cultured in a mixture of 98% glucose free RPMI 1640 supplemented with 10% dialyzed fetal calf serum and 2% of regular RPMI 1640 medium containing 10% fetal calf serum (final glucose concentration, 0.28 mM). The medium was supplemented with sodium pyruvate (0.11 mg/ml), penicillin (100 units/ml), and streptomycin (0.1 mg/ml). The cells were cultured for 24 h in the presence of [<sup>3</sup>H] glucosamine or [<sup>3</sup>H] galactose [DuPont - New England Nuclear] at a concentration of 20  $\mu$ Ci/ml.

**Immunoprecipitation and sodium dodecyl sulfate-polyacrylamide gel electrophoresis** - Radiolabeled cells were lysed in PBS (6.7 mM potassium phosphate buffer, pH 7.4, 150 mM NaCl, 0.002% NaN<sub>3</sub>) containing 1% Nonidet P-40 and protease inhibitors as described.<sup>19</sup> Lamp-2 was immunoprecipitated from the cell lysates by the addition of rabbit anti-human lamp-2 antibodies followed by the addition of formalin-fixed *Staphylococcus aureus* (Pansorbin, Calbiochem) as described.<sup>19</sup> The supernatant from this immunoprecipitation was then subjected to immunoprecipitation using anti-human lamp-1 antibodies.<sup>20</sup> Aliquots of radiolabeled samples were subjected to sodium dodecyl sulfate-polyacrylamide gel electrophoresis on a 8% polyacrylamide gel according to Laemmli,<sup>21</sup> followed by fluorography using Enlightning (DuPont - New England Nuclear) (Fig. 1).

**Lectin Affinity Chromatography of Glycopeptides** - Glycopeptides were released by Pronase digestion of immunoprecipitates and fractionated by serial lectin affinity chromatography as described<sup>19</sup> except for Jacalin-Agarose chromatography. Briefly, the glycopeptides were first applied to a column of concanavalin A-Sepharose, and highly branched complex-type *N*-glycan glycopeptides and *O*-glycan containing glycopeptides (Fraction I) were separated from biantennary asparagine-linked glycopeptides (Fraction II) and high-mannose glycopeptides (Fraction III). Fraction I was then applied to *Datura stramonium* agglutinin (DSA)-Agarose. The glycopeptides bound to this column would contain *N*-glycans with poly-*N*-acetylglucosamine repeats and/or R→GlcNAc $\beta$ 1→6 (R→GlcNAc $\beta$ 1→2)Man→ branchings (Fraction I B). The glycopeptides unbound to this column were further applied to a column (0.7 x 10.5 cm) of Jacalin-Agarose (Vector Laboratories, Burlingame,

CA). The column was equilibrated with 100 mM Tris-HCl, pH 7.5 containing 0.5 M NaCl,<sup>22</sup> and the glycopeptides dissolved in the same buffer were applied to the column. After washing with the same buffer, the glycopeptides bound to Jacalin-Agarose were eluted with 0.1 M melibiose in the same buffer (Aldrich).<sup>17,22</sup> In order to ensure the complete binding of glycopeptides, the flow rate was maintained at 4 ml/h throughout the procedure. These glycopeptides were desalted by gel filtration on a column of Sephadex G-25 eluted with water containing 7% propanol.

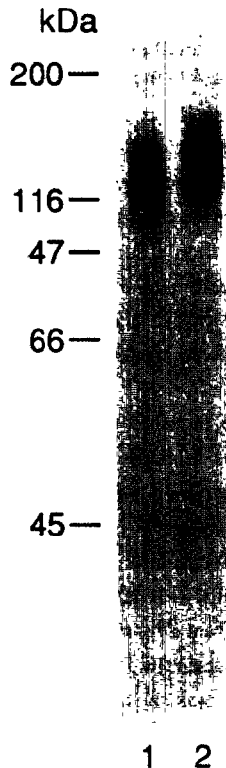


Fig. 1 Sodium dodecylsulfate-polyacrylamide gel electrophoresis of lamp-1 and lamp-2 from HL-60 cells. HL-60 cells were labeled with [<sup>3</sup>H] glucosamine and lamp-1 and lamp-2 were sequentially immunoprecipitated (lanes 1 and 2). They were then subjected to sodium dodecylsulfate-polyacrylamide gel electrophoresis followed by fluorography. The molecular weights of standard proteins migrated are indicated at the left.

In order to obtain the whole hinge-region as one glycopeptide, the immunoprecipitates of lamp-1 and lamp-2 were eluted by boiling in 0.1 % SDS for 5 min. The supernatants obtained after brief centrifugation were subjected to reduction with dithiothreitol followed by alkylation with monoiodoacetic acid.<sup>20</sup> To the reduced and alkylated sample was then added 3 volumes of acetone, and the suspension was briefly centrifuged. The precipitates, dissolved in 0.1 M ammonium bicarbonate buffer, pH 8.0, were digested at 37°C with 80 µg

of trypsin for lamp-1 or 40  $\mu$ g of trypsin followed by 4  $\mu$ g of V8 protease for lamp-2.<sup>20</sup> Total incubation time was 28 h for lamp-1 or 44 h for lamp-2. The digests were applied to a short column of Sephadex G-25 for desalting before being subjected to serial lectin affinity chromatography.

Alkaline Borohydride Treatment of Glycopeptides - In order to release *O*-glycans from glycopeptides, the glycopeptides were incubated in 0.05 M NaOH - 1 M NaBH<sub>4</sub> at 45°C for 48 h. After this treatment, the sample was treated with methanol containing 1 M acetic acid and methylborate was removed under a flow of nitrogen in a water bath at 37°C. After repeating this treatment one more time, methanol was added to the sample and then removed as described above. The dried sample was either immediately subjected to Sephadex G-15 gel filtration for desalting or frozen at -20°C until desalting.

Fractionation of *O*-glycans and *N*-glycans by Gel Filtration - *O*-Glycans were fractionated by gel filtration using a column (1.0 x 105 cm) of Bio-Gel P-4 (200 to 400 mesh) equilibrated with 0.1 M NH<sub>4</sub>HCO<sub>3</sub>. *N*-glycans were fractionated by gel filtration using a column (1.0 x 105 cm) of Sephadex G-50 (Superfine) equilibrated with 0.1 M NH<sub>4</sub>HCO<sub>3</sub>. Fractions of 1 ml were collected at a flow rate of 5 ml/h.

Glycosidase Treatment - Radiolabeled *O*-glycans were digested with 90 munits of endo- $\beta$ -galactosidase in 100  $\mu$ l of 50 mM sodium acetate buffer, pH 5.8 under the conditions described previously.<sup>23</sup> Endo- $\beta$ -galactosidase was purified from *Escherichia freundii*<sup>23</sup> and kindly donated by Dr. Michiko N. Fukuda of our institute. Radiolabeled glycopeptides or *O*-glycans were digested with 40 munits of Newcastle disease virus neuraminidase in 80  $\mu$ l of 50 mM cacodylate buffer pH6.5 at 37°C for 24 h.<sup>17</sup> It has been shown that Newcastle disease virus neuraminidase can cleave the NeuNAc $\alpha$ 2 $\rightarrow$ 3 linkage but barely cleaves other linkages or not at all.<sup>24</sup>

The glycopeptides and *O*-glycans were digested with a mixture of 15 munits of diplococcal  $\beta$ -galactosidase and 20 munits of  $\beta$ -*N*-acetylglucosaminidase in 110  $\mu$ l of 50 mM cacodylate buffer, pH 6.0. If necessary, 50 mM of 2,3-dehydro-2-deoxy-*N*-acetylneuraminic acid (Boehringer-Mannheim) was added as an inhibitor for potentially contaminating neuraminidases.<sup>25</sup> If necessary, 100 mM of L-fucose (Sigma) was also added to inhibit potentially contaminating  $\alpha$ -L-fucosidase. As shown previously, diplococcal  $\beta$ -galactosidase can cleave Gal $\beta$ 1 $\rightarrow$ 4GlcNAc linkage but not Gal $\beta$ 1 $\rightarrow$ 3GlcNAc linkage.<sup>26</sup> Glycopeptides and *O*-glycans were digested with 20 munits of  $\alpha$ 1 $\rightarrow$ 3/4 fucosidase from *Streptomyces sp.* 142,<sup>27</sup> together with the same amount of diplococcal  $\beta$ -galactosidase and  $\beta$ -*N*-acetylglucosaminidase as described above, in 90  $\mu$ l of 25 mM citrate phosphate buffer, pH 6.0. This  $\alpha$ -L-fucosidase was shown to cleave Gal $\beta$ 1 $\rightarrow$ 4(Fuc  $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ R and Gal $\beta$ 1 $\rightarrow$ 3(Fuc  $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ R but not NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ R or NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 3(Fuc $\alpha$ 1 $\rightarrow$ 4)GlcNAc $\beta$ 1 $\rightarrow$ R under these conditions.<sup>17,27</sup> All these digestions were carried out at 37°C for 48 h. Newcastle disease virus neuraminidase was prepared in our laboratory whereas diplococcal  $\beta$ -galactosidase and  $\beta$ -*N*-acetylglucosaminidase were purchased from Boehringer-Mannheim. *Streptomyces*  $\alpha$ 1 $\rightarrow$ 3/4 fucosidase was purchased from Takara (Berkeley, CA). Standard oligosaccharides for *O*-glycans and digestion products after endo- $\beta$ -galactosidase treatment were obtained from leukosialin and lamp-1 and lamp-2 as described previously.<sup>19,28,29</sup>

## RESULTS

Isolation of *O*-Glycans from Lamp-1 and Lamp-2 - Lamp-1 and lamp-2 are extremely enriched with *N*-glycans and less than 10% of the total carbohydrate is due to *O*-glycans. In order to isolate selectively *O*-glycans, glycopeptides were subjected to serial lectin affinity chromatography as shown in Fig. 2. In this

chromatography, the glycopeptides containing *O*-glycans are not bound to ConA-Sepharose nor DSA-Agarose, but should bind to Jacalin-Agarose.

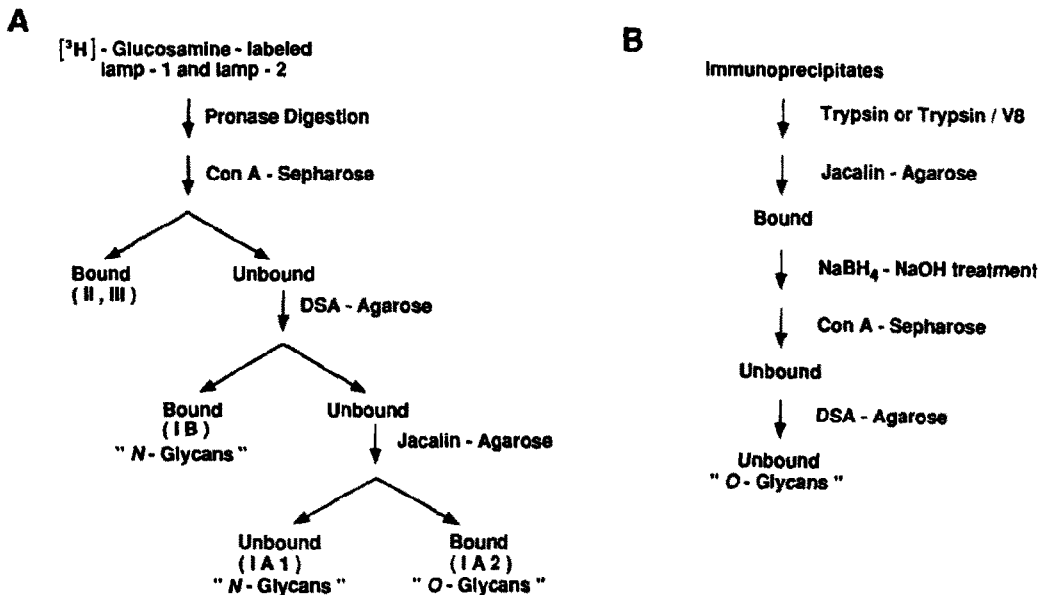


Fig. 2 Experimental procedures used for the isolation of glycopeptides containing *N*-glycans and *O*-glycans.

A) The samples shown in Fig. 1 were digested by Pronase and the radio-labeled glycopeptides were sequentially applied to ConA-Sepharose, DSA-Agarose, followed by Jacalin-Agarose. Fraction IB and Fraction IA1 are glycopeptides containing *N*-glycans whereas Fraction IA2 contain glycopeptides containing *O*-glycans.

B) The samples shown in Fig. 1 were digested by trypsin (for lamp-1) or trypsin followed by V8 protease (for lamp-2) and the radiolabeled glycopeptides were applied to a column of Jacalin-Agarose. The fraction bound and eluted from Jacalin-Agarose are glycopeptides containing *O*-glycans. Since this fraction was found to contain contaminating *N*-glycans, they were removed by ConA-Sepharose followed by DSA-Agarose after alkaline borohydride treatment. This fraction, after ConA- and DSA-column purification, still contains a small amount of DSA-unbound *N*-glycans in addition to *O*-glycans (see the text).

First, glycopeptides were prepared by pronase digestion of [<sup>3</sup>H]-glucosamine labeled lamp-1 and lamp-2 of HL-60 cells and subjected to serial lectin affinity chromatography as shown in Fig. 2A. The glycopeptides bound to Jacalin-Agarose (5-6% of the total radioactivity) were then treated with alkaline borohydride, and the released *O*-glycans were subjected to Bio-Gel P-4 gel filtration. As shown in Fig. 3A and B, those *O*-glycans were separated into 8 fractions, and the structures of oligosaccharides at fractions 3 to 9 were determined as follows.

Oligosaccharide 3 produced the tetrasaccharide, Galβ1→3(Galβ1→4GlcNAcβ1→6)GalNAcOH and two sialic acid residues after treatment with Newcastle disease virus neuraminidase, indicating that oligosaccharide 3 is disialosylhexasaccharide, NeuNAcα2→3Galβ1→3(NeuNAcα2→3Galβ1→4GlcNAcβ1→6)GalNAcOH. As a minor component (less than 10%) of the total radioactivity, a radioactive peak corresponding to

Gal $\beta$ 1 $\rightarrow$ 3[Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 6]GalNAcOH was produced, indicating the presence of NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 3[NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 6]GalNAcOH. Oligosaccharide 5 yielded one sialic acid and the tetrasaccharide, after treatment with Newcastle disease virus neuraminidase. Further treatment with  $\beta$ -galactosidase and  $\beta$ -*N*-acetylglucosaminidase yielded Gal $\beta$ 1 $\rightarrow$ 3GalNAcOH, indicating oligosaccharide 5 being monosialosyl pentasaccharide. In order to determine the position of sialic acid substitution in galactose residues, oligosaccharide 5 from lamp-1 (Fig. 3C) was digested with a mixture of  $\beta$ -galactosidase and  $\beta$ -*N*-acetylglucosaminidase from jack bean. After this treatment, 48% of the radioactivity was converted to GlcNAc and monosialosyltrisaccharide. However, 52% of radioactivity remained intact. These results indicate that oligosaccharide 5 is a mixture of NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 3(Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 6)GalNAcOH and Gal $\beta$ 1 $\rightarrow$ 3(NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 6)GalNAcOH in a ratio of 48:52. Similarly, the structures of oligosaccharides 6-9 were determined as shown in Table I.

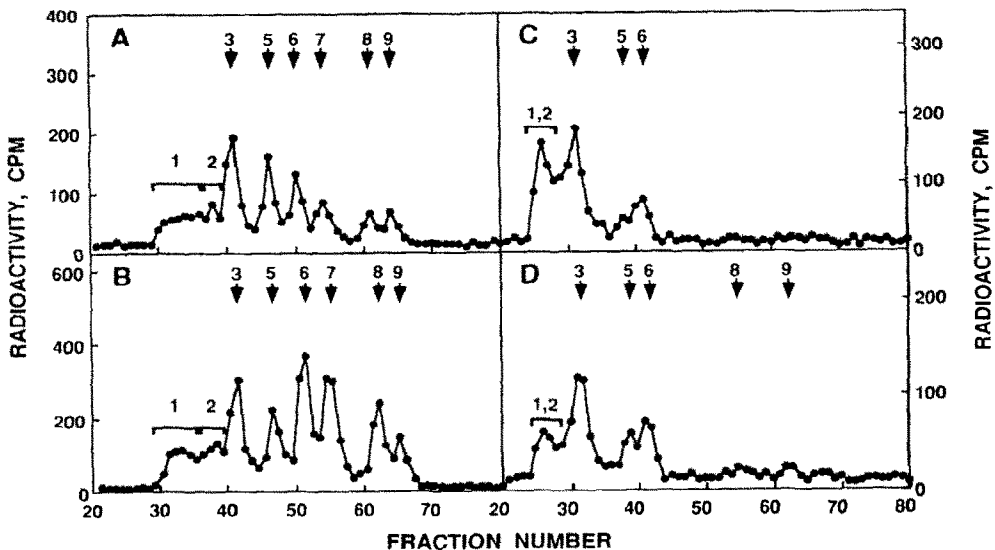


Fig. 3 Bio-Gel P-4 gel filtration of *O*-glycans released from glycopeptides bound to Jacalin-Agarose.

The lamp-1 and lamp-2 glycopeptides bound to Jacalin-Agarose were treated with alkaline borohydride and subjected to Bio-Gel P-4 gel filtration. **A, B**, *O*-glycans released from Pronase digested glycopeptides of lamp-1 (A) and lamp-2 (B). **C, D**, *O*-glycans released from trypsin digested lamp-1 glycopeptides (C) and from trypsin-V8 protease digested lamp-2 glycopeptides (D). Peak 1 and peak 2 represent *O*-glycans containing poly-*N*-acetylglucosamine in A and B. In addition, DSA-unbound *N*-glycans are present in peaks 1 and 2 of C and D. The elution positions of standard oligosaccharides are as follows: 3, NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 3 (NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 6) GalNAcOH; 5, NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 3 (Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 6) GalNAcOH plus Gal $\beta$ 1 $\rightarrow$ 3 (NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 6) GalNAcOH; 6, NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 3GalNAcOH; 7, Gal $\beta$ 1 $\rightarrow$ 3(Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 6)GalNAcOH; 8, Gal $\beta$ 1 $\rightarrow$ 3GalNAcOH; 9, GalNAcOH.

TABLE I

Structure and relative molar ratios of the *O*-linked oligosaccharides isolated from lamp-1 and lamp-2 and leukosialin from HL-60 cells.

Structure	Lamp-1	Lamp-2	Leukosialin <sup>a</sup>
GalNAcOH	3.4%	2.1%	13.3%
Galβ1→3GalNAcOH	5.1%	5.1%	2.9%
NeuNAcα2→3Galβ1→3GalNAcOH	22.7%	30.0%	7.1%
NeuNAcα2 ↓ 6 NeuNAcα2→3Galβ1→3GalNAcOH	.b	-	-
Galβ1→4GlcNAcβ1 ↓ 6 Galβ1→3GalNAcOH	3.0%	2.7%	4.8%
NeuNAcα2→3Galβ1→4GlcNAcβ1 ↓ 6 Galβ1→3GalNAcOH	4.5%		
Galβ1→4GlcNAcβ1 ↓ 6 NeuNAcα2→3Galβ1→3GalNAcOH		14.3%	8.6%
NeuNAcα2→3Galβ1→3GalNAcOH	4.1%		
NeuNAcα2→3Galβ1→4GlcNAcβ1 ↓ 6 NeuNAcα2→3Galβ1→3GalNAcOH	47.8%	37.3%	60.6%

HL-60, lamp-1, and lamp-2 contain approximately 21% and 12% of *O*-glycans as poly-*N*-acetylglucosaminyl *O*-glycans, respectively. <sup>a</sup>The data on leukosialin is in reference 17. <sup>b</sup>Less than 1%.

As shown previously, *O*-glycans are exclusively attached to the hinge-region of lamp-1 and lamp-2.<sup>20</sup> We therefore digested lamp molecules with trypsin (for lamp-1) or trypsin followed by V8 protease (for lamp-2). These digestion conditions allow us to obtain the whole hinge-region in one glycopeptide and yet excluding the *N*-glycosylation site (see Fig. 3 in reference 20). The glycopeptides obtained were then applied to a column of Jacalin-Agarose and the glycopeptides bound to this column were subjected to alkaline borohydride treatment. The amount of the glycopeptides bound to the Jacalin-column was about 50% more than the glycopeptides prepared after pronase. In order to remove *N*-glycans, the released glycans were applied to Con A-Sepharose and DSA-Agarose (Fig. 2B). *O*-Glycans unbound to these two lectins were then subjected to Bio-

Gel P-4 gel filtration. As shown in Fig. 3C and D, the oligosaccharides derived from the whole hinge-region are much more enriched in the disialosyl hexasaccharide (peak 3). The ratio of oligosaccharides 3-9 obtained from lamp molecules is almost the same as that obtained in leukosialin (Table I). These results indicate that the fully sialylated hexasaccharide was recovered in Jacalin-bound fraction because the glycopeptides contained small oligosaccharides that can be bound to Jacalin. The Jacalin-unbound fraction was desialyzed and again applied to the Jacalin column, but no additional glycopeptides were found, indicating that *O*-glycans were fully recovered in Fig. 3C and D.

Glycosidase Digestion of Poly-*N*-Acetylglucosaminyl *O*-Glycans - Peaks 1 and 2 obtained after Bio-Gel P-4 gel filtration (Fig. 3) were found to contain sialyl Le<sup>x</sup> structures in poly-*N*-acetylglucosaminyl *O*-glycans, as summarized below.

- (1) Peak 2 in Fig. 3A and B is larger than the disialosylhexasaccharide by one *N*-acetylglucosaminyl unit. This peak 2 eluted at the same position of the oligosaccharide NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4( $\pm$ Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 6(NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 3)GalNAcOH, which was isolated from leukosialin.<sup>17</sup>
- (2) Both the peaks 1 and 2 were susceptible to endo- $\beta$ -galactosidase (Fig. 4).

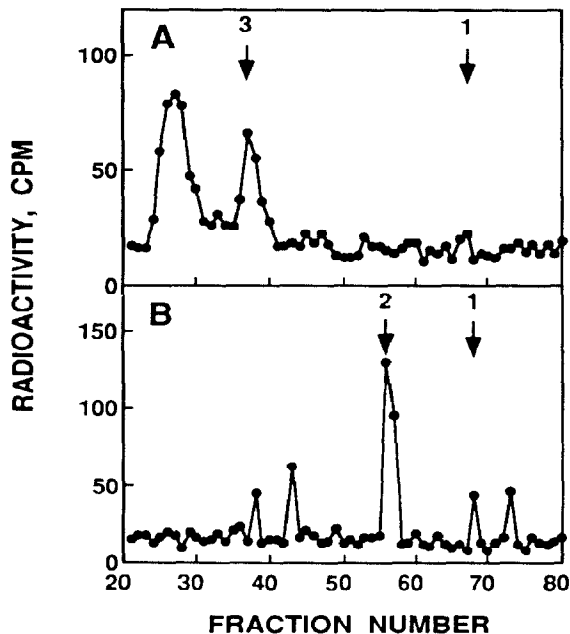


Fig. 4 Bio-Gel P-4 gel filtration of lamp-2 peak 1 after endo- $\beta$ -galactosidase digestion.

Peak 1 in Fig. 3B was digested with endo- $\beta$ -galactosidase before (A) and after desialylation (B). The elution positions of standard oligosaccharide are: 1, GlcNAc $\beta$ 1 $\rightarrow$ 3Gal; 2, Gal $\beta$ 1 $\rightarrow$ 3(GlcNAc $\beta$ 1 $\rightarrow$ 6)GalNAcOH plus Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 3Gal; 4, NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3Gal. Similar results were obtained on the lamp-1 sample.



- (3) Both peak 1 and peak 2 are exclusively composed of sialylated oligosaccharides (Fig. 5A). Without prior treatment of sialidase, no release of Gal or GlcNAc was detected (data not shown). Peak 1 plus peak 2 yielded Gal $\beta$ 1 $\rightarrow$ 3GalNAcOH after digestion of neuraminidase,  $\beta$ -galactosidase and  $\beta$ -*N*-acetylglucosaminidase (Fig. 5B). This amount was increased when the oligosaccharides were digested with  $\alpha$ 1 $\rightarrow$ 3/4 fucosidase in addition to  $\beta$ -galactosidase and  $\beta$ -*N*-acetylglucosaminidase (Fig. 5C).

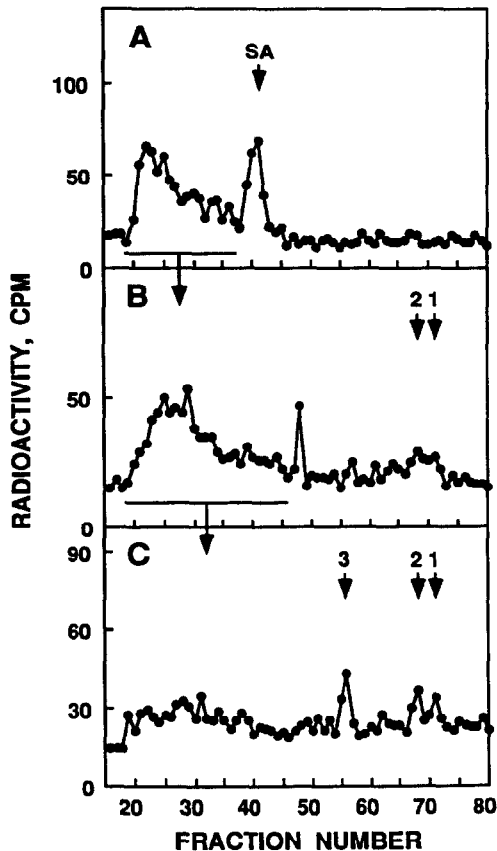


Fig. 5 Sequential glycosidase treatment of peak 1 plus peak 2 from lamp-1 and lamp-2.

Peak-1 plus peak 2 in Fig. 3C were digested with neuraminidase (A) then  $\beta$ -galactosidase plus  $\beta$ -*N*-acetylglucosaminidase (B) followed by  $\beta$ -galactosidase,  $\beta$ -*N*-acetylglucosaminidase plus  $\alpha$ -fucosidase (C). The digest was subjected to Bio-Gel P-4 gel filtration. The elution position of the standard oligosaccharides are 1, GlcNAc; 2, Gal $\beta$ 1 $\rightarrow$ 3GalNAcOH; 3, (Man) $_3$ Fuc(GlcNAc) $_2$ . (Man) $_3$ Fuc(GlcNAc) $_2$  was produced because alkaline borohydride treatment cleaved asparitylglycosylamine linkage.<sup>35</sup>



were treated with alkaline borohydride. The results indicate only a very small amount of the radioactivity was due to *O*-glycans, shifting into low molecular weight after the treatment (Figs. 6A and 7A).

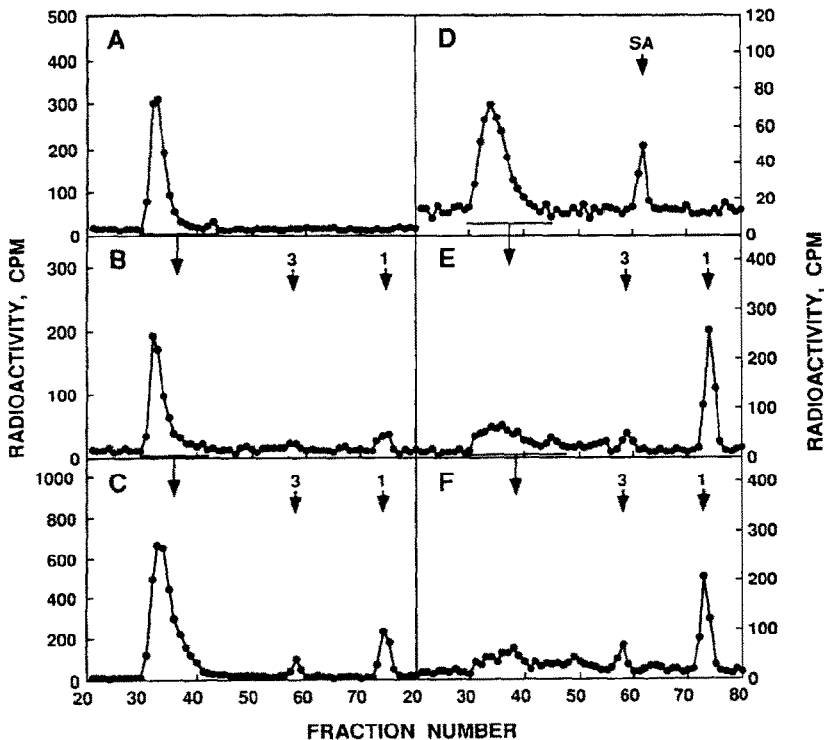


Fig. 6 Sequential glycosidase digestion of Jacalin unbound glycopeptides, IA1 obtained from lamp-1.

Glycopeptides that were not bound to DSA-Agarose or Jacalin-Agarose (IA1 in Fig. 2A) were subjected to Bio-Gel P-4 gel filtration after alkaline borohydride treatment (A). The glycopeptides after alkaline borohydride treatment were then digested with  $\beta$ -galactosidase plus  $\beta$ -*N*-acetylglucosaminidase (B) followed by  $\beta$ -galactosidase,  $\beta$ -*N*-acetylglucosaminidase plus  $\alpha$ -fucosidase (C). In parallel, the same glycopeptides after alkaline borohydride treatment were digested with neuraminidase (D), then by  $\beta$ -galactosidase plus  $\beta$ -*N*-acetylglucosaminidase (E) followed by  $\beta$ -galactosidase,  $\beta$ -*N*-acetylglucosaminidase plus  $\alpha$ -fucosidase (F). The bars indicate the fractions pooled for further digestion. The elution positions of standard oligosaccharides are: 1, GlcNAc; SA, sialic acid; 3, (Man)<sub>3</sub>Fuc(GlcNAc)<sub>2</sub>. (Man)<sub>3</sub>Fuc(GlcNAc)<sub>2</sub> was obtained because *N*-glycans were partially released from glycopeptides by alkaline borohydride treatment.<sup>35</sup>

The glycopeptides remaining after alkaline borohydride treatment were then subjected to sequential exoglycosidase digestion. Figure 6 illustrates that only a small portion of *N*-glycans in lamp-1 IA1 are susceptible to  $\beta$ -galactosidase plus  $\beta$ -*N*-acetylglucosaminidase prior to neuraminidase treatment. By comparing Fig. 6B and C, it can be estimated that 9.8% of *N*-acetylglucosamine is Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3 and 11.2% is

Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 3. Similar analysis was made after neuraminidase treatment (Fig. 6D and E), and it can be estimated that Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3 constitutes 22.8% of *N*-acetylglucosamine whereas Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 3 constitutes 24.8% in the desialylated sample. The difference thus obtained on the results before and after desialylation should yield the amount of sialylated forms. Thus lamp-1 IA-1 contains 13.0% of NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3 and 13.6% of NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 3.

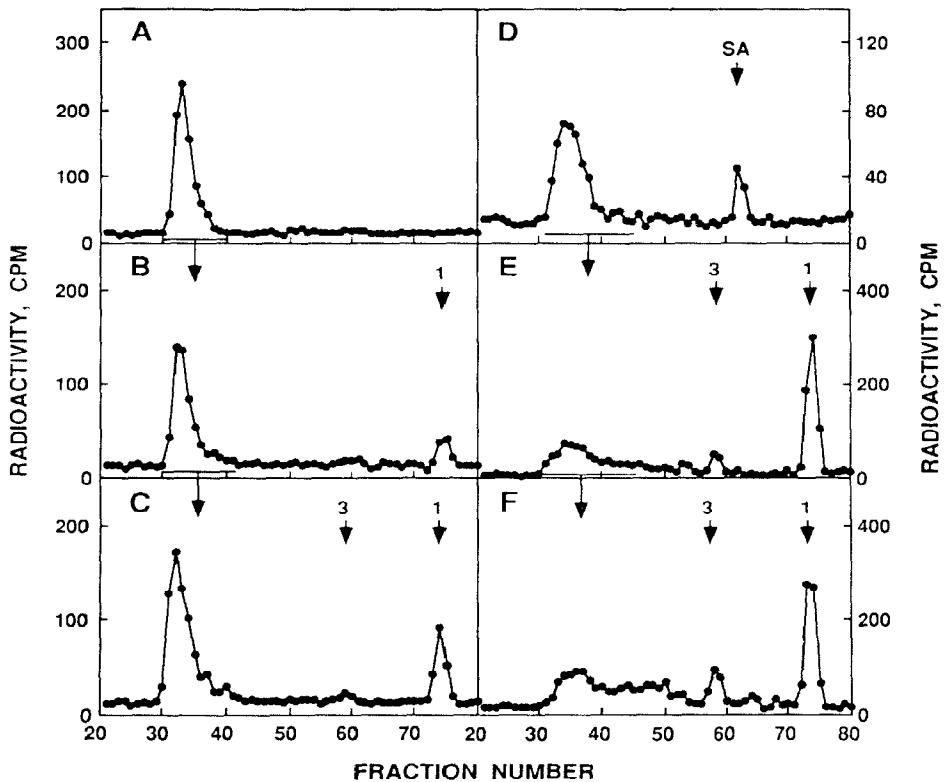


Fig. 7 Sequential glycosidase digestion of DSA-bound fraction (IB) from lamp-1.

Lamp-1 glycopeptides that were bound to DSA-Agarose (IB in Fig. 2A) were subjected to Bio-Gel P-4 gel filtration after alkaline borohydride treatment (A). The glycopeptides after alkaline borohydride treatment were then digested with  $\beta$ -galactosidase plus  $\beta$ -*N*-acetylglucosaminidase (B) followed by  $\beta$ -galactosidase,  $\beta$ -*N*-acetylglucosaminidase plus  $\alpha$ -fucosidase (C). In parallel, the glycopeptides after alkaline borohydride treatment were digested with neuraminidase (D), then  $\beta$ -galactosidase plus  $\beta$ -*N*-acetylglucosaminidase (E) followed by  $\beta$ -galactosidase,  $\beta$ -*N*-acetylglucosaminidase plus  $\alpha$ -fucosidase (F). The elution positions of standard oligosaccharides are as shown in Fig. 6. The bar indicates the fractions pooled for further digestion.

We have previously shown<sup>19</sup> that IA1 fraction contains a total of 6.5 moles of *N*-acetylglucosamine. It was also determined that the IA1 fraction contains 4.2 moles of *N*-glycans/mole of lamp-1, judging from the recovery of glycopeptides in each fraction and the total number of *N*-glycans.<sup>19</sup> Based on this data, it can be calculated that the IA1 fraction in the present study represents the total of 27.3 moles of *N*-acetylglucosamine residues ( $6.5 \times 4.2 = 27.3$ ). Based on this total amount of *N*-acetylglucosamine and the amount of each structure determined above, the amount of Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3, Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 3, NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3, and NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\beta$ 1 $\rightarrow$ 3, can be calculated.

Similar calculations can be made on DSA-bound fraction (IB) obtained from lamp-1, based on the yield of *N*-acetylglucosamine after sequential glycosidase digestion (Fig. 7). The amount of each structure in lamp-1 was obtained by summation of those present in IA1 and IB and summarized in Table II. Based on similar experiments, the amount of each structure can be calculated on lamp-2 (Table II).

**TABLE II**  
The amount of *N*-acetylglucosamines with Le<sup>x</sup> or sialyl Le<sup>x</sup> terminus in lamp-1 and lamp-2 of HL-60 cells.

Structure	Lamp-1		Lamp-2		Leukosialin <sup>a</sup>
	<i>N</i> -glycans	<i>O</i> -glycans	<i>N</i> -glycans	<i>O</i> -glycans	<i>O</i> -glycans
	moles/mole of glycoproteins				
(Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3) <sub>n</sub>	12.3 <sup>b</sup>	0	11.0	0	0
(Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3) <sub>3</sub> ↑ $\alpha$ 1 Fuc	13.4	0	11.8	0	0
NeuNAc $\alpha$ 2 $\rightarrow$ 3(Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3) <sub>n</sub>	26.6	0.6	31.4	0.3	0.7
NeuNAc $\alpha$ 2 $\rightarrow$ 3(Gal $\beta$ 1 $\rightarrow$ 4GlcNAc $\beta$ 1 $\rightarrow$ 3) <sub>3</sub> ↑ $\alpha$ 1 Fuc	4.7	1.1	3.2	0.7	0.5

a. The data on leukosialin is in reference 17. b. The values are based on the numbers of *N*-glycans and *O*-glycans attached to each molecule and obtained by calculation of the yield of each structure.

## DISCUSSION

The present studies demonstrated that *O*-glycans present in lysosomal membrane glycoproteins, lamp-1 and lamp-2, contain poly-*N*-acetylglucosaminyl repeats when the proteins were isolated from HL-60 cells. The studies also demonstrated that most of these poly-*N*-acetylglucosaminyl *O*-glycans are terminated with the sialyl Le<sup>x</sup> structure, NeuNAc $\alpha$ 2 $\rightarrow$ 3Gal $\beta$ 1 $\rightarrow$ 4(Fuc $\alpha$ 1 $\rightarrow$ 3)GlcNAc $\rightarrow$ R. The results also show that the amount of sialyl Le<sup>x</sup> in *O*-glycans is significant, considering that only a fraction of *N*-glycans contain the sialyl Le<sup>x</sup> terminus (Table II).

Previously, we have shown only 1.5% of *O*-glycans attached to leukosialin contain poly-*N*-acetylglucosaminyl extension and 40% of those poly-*N*-acetylglucosaminyl *O*-glycans contain sialyl Le<sup>x</sup> terminus. Since leukosialin contains approximately 80 *O*-glycans per molecule, these results indicate that leukosialin contains, on average, 0.5 mole of sialyl Le<sup>x</sup> terminus. In contrast, despite the fact that only 5 to 8 *O*-glycans are present in lamp molecules, *O*-glycans in lamp molecules contain slightly more sialyl Le<sup>x</sup> terminus per molecule than leukosialin does. This is mostly because *O*-glycans attached to lamp molecules contain about fifteen times more poly-*N*-acetylglucosaminyl repeats than leukosialin. Lamp molecules were isolated in our laboratory as the major sialoglycoproteins containing *N*-acetylglucosamine repeats in *N*-glycans.<sup>18</sup> The present study thus demonstrated that both *N*- and *O*-glycans in lamp molecules are modified by poly-*N*-acetylglucosamine extension more efficiently than those attached to leukosialin. These results support the previous findings that poly-*N*-acetylglucosamine formation is somehow intrinsic to the nature of glycoproteins.<sup>1,2</sup> Poly-*N*-acetylglucosamines are formed when  $\beta$ -1,3-*N*-acetylglucosaminyltransferase and  $\beta$ -1,4-galactosyltransferase alternatively add *N*-acetylglucosamine and galactose to *N*-acetylglucosamine, the precursor (Fig. 8). We found that glycoproteins containing a significant amount of poly-*N*-acetylglucosamine go through the Golgi complex slowly so that the enzymes adding *N*-acetylglucosamine repeats can act on those glycoproteins.<sup>30</sup> It is also possible that lamp molecules present the precursor carbohydrate in a unique way so that *N*-acetylglucosamine repeats are efficiently formed by alternate additions of *N*-acetylglucosamine and galactose.

It has been shown previously that leukosialin from HL-60 cells contains sialyl Le<sup>x</sup> terminus but not leukosialin from K562 cells.<sup>17</sup> *O*-Glycans from HL-60 cells contain those based on core 2 branchings which are formed by core 2  $\beta$ -1,6-*N*-acetylglucosaminyltransferase and those branched oligosaccharides can be modified to acquire poly-*N*-acetylglucosaminyl repeats when  $\beta$ -1,3-*N*-acetylglucosaminyltransferase is also present (Figure 8). Moreover, those poly-*N*-acetylglucosaminyl extensions apparently acquire sialyl Le<sup>x</sup> terminus much more efficiently than one *N*-acetylglucosaminyl branch attached to Gal $\beta$ 1 $\rightarrow$ 3GalNAc. This is probably because  $\alpha$ -1,3-fucosyltransferase adds more efficiently a fucose to sialyl *N*-acetylglucosamine when the terminal structure is present at the end of poly-*N*-acetylglucosamine extension.

In order to form sialyl Le<sup>x</sup> structure in *O*-glycans, the formation of core 2 branch is thus a prerequisite. In fact, we have shown recently that the expression of cloned core 2  $\beta$ -1,6-*N*-acetylglucosaminyltransferase made Chinese hamster ovary cells express poly-*N*-acetylglucosamine extensions in *O*-glycans.<sup>31</sup> The parent CHO cells lacking this enzyme were found not to contain poly-*N*-acetylglucosaminyl *O*-glycans. These results clearly indicate that core 2  $\beta$ -1,6-*N*-acetylglucosaminyltransferase is the key enzyme for sialyl Le<sup>x</sup> terminus in *O*-glycans (see Figure 8).

In relation to this, it is noteworthy that human T-cell activation was found to be associated with the appearance of core 2-based oligosaccharides.<sup>16</sup> It was found later that sialyl Le<sup>x</sup> structure appears on T cells

when they are activated.<sup>32</sup> These results strongly suggest that core 2 formation is also the key step for sialyl Le<sup>x</sup> formation in activated T-lymphocytes.

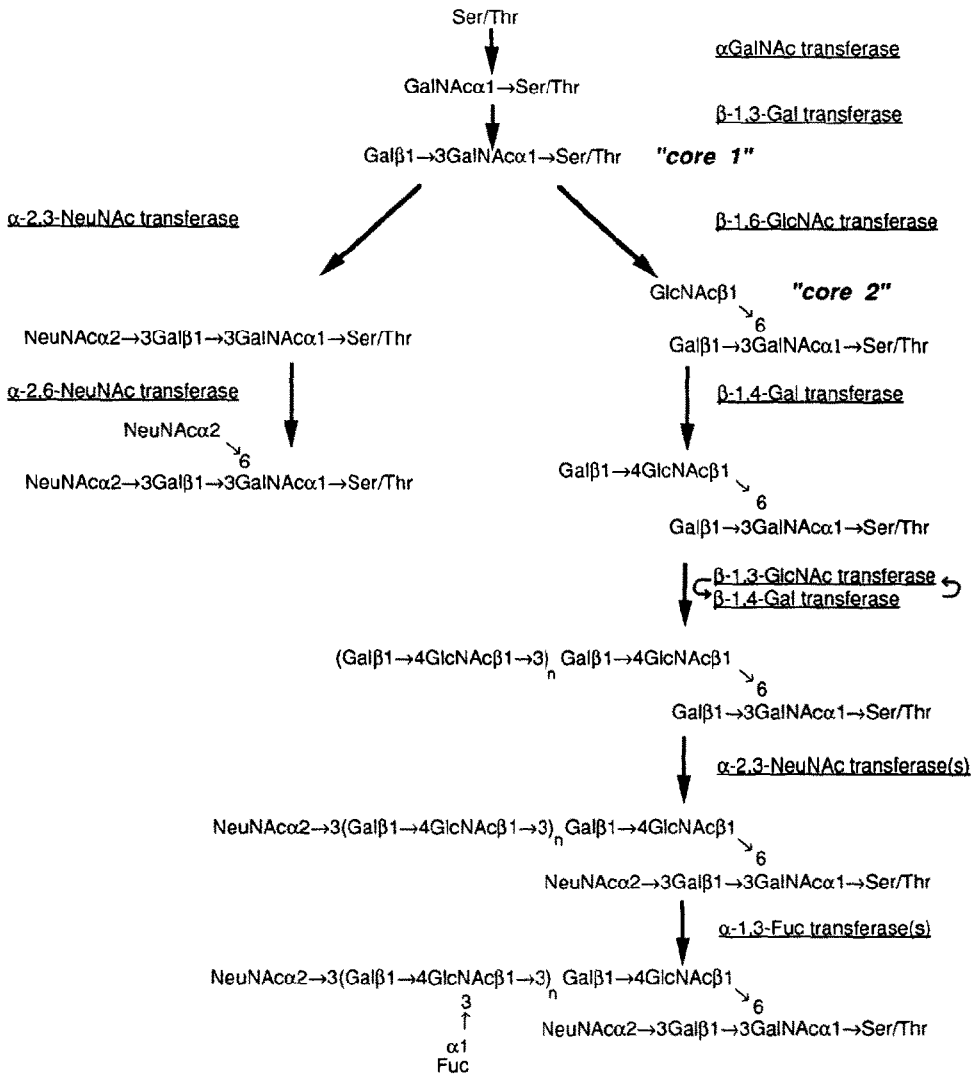


Fig. 8 The proposed biosynthetic pathways of O-glycans and poly-N-acetylglucosaminyl O-glycans. It has been shown that the tetrasaccharide (bottom left) is formed by sequential addition of  $\alpha 2 \rightarrow 3$  sialyltransferase followed by  $\alpha 2 \rightarrow 6$  sialyltransferase. When  $\beta 1 \rightarrow 6$  N-acetylglucosaminyltransferase, core 2 enzyme, is present, the core 2 branch is formed. To this core 2 branch, poly-N-acetylglucosaminyl extension can be added. Poly-N-acetylglucosaminyl chain can be further modified by  $\alpha 1 \rightarrow 3$  fucosyltransferase, forming sialyl Le<sup>x</sup> terminus (based on Refs. 6, 12, 17, 26 and 31).

It has been demonstrated in several laboratories that the sialyl Le<sup>x</sup> structure is the ligand for E- and P-selectins.<sup>6-9</sup> Recent studies revealed that sialyl Le<sup>x</sup> structure is more efficiently recognized by selectin when they are present in mucin-type glycoproteins.<sup>33,34</sup> The results strongly suggest that presentation of ligands in *O*-glycans leads into higher affinity to selectins because they are present as a cluster. Since the formation of sialyl Le<sup>x</sup> in *O*-glycans is determined by core 2 branchings, further studies are significant to understand how the expression of core 2 forming  $\beta$ -1,6-*N*-acetylglucosaminyltransferase is regulated.

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