Lectin-carbohydrate recognition between the selectins and their ligands are among the earliest events in leukocyte recirculation, leukocyte recruitment into inflamed areas, and abnormal egress of leukocytes in diseases. Previously, we have described a dimeric sialoglycoprotein from myeloid cells with subunits of molecular mass = 120 kDa, which is selectively recognized by P-selectin (Moore, K. L., Stults, N. L., Diaz, S. M., Smith, D. F., Cummings, R. D., Varki, A., and McEver, R. P. (1992) J. Cell Biol. 188, 445-456).

Here, we demonstrate that this P-selectin ligand carries 2-3-linked sialic acids and the sialyl-Lewis\(^+\) (SLe\(^+\)) tetrasaccharide motif. This glycoprotein contains <1% of the total membrane-bound sialic acids and a very small fraction of the total SLe\(^+\) on neutrophil membranes. In spite of a relative stability to sialidase digestion, the predominant form of sialic acid on the ligand is N-acetyllactosaminic acid. Selective periodate oxidation of the side chain of sialic acids does not affect P-selectin binding and allows the introduction of tritium label into the truncated sialic acids. \(\beta\)-Elimination with alkaline borohydride releases labeled O-linked oligosaccharides both from the labeled neutrophil ligand and from the ligand purified from HL-60 cells metabolically labeled with \[^{[\text{3}H]\text{glucosamine}}\]. The ligand from both neutrophils and HL-60 cells is also susceptible to cleavage by the enzyme O-sialoglycoprotease from *Pasteurella hemolytica*. Analysis of the specificity of this enzyme suggests that the P-selectin ligand carries large numbers of closely spaced sialylated O-linked oligosaccharides. O-Sialoglycoprotease abolishes both direct binding of P-selectin to HL-60 cells and the adhesion of HL-60 cells to immobilized P-selectin, without significantly decreasing overall cell surface SLe\(^+\) expression. This indicates that the 120-kDa ligand may be the major determinant of P-selectin/myeloid cell interaction in vivo. Finally, based on the current and previous data, we hypothesize that the high affinity recognition site(s) of this P-selectin ligand may be derived from a "clustered saccharide patch" of sialylated fucosylated O-linked oligosaccharide sequences.

The selectins are a family of calcium-dependent mammalian lectins that initiate interactions between blood cells and endothelium (1-7). A large body of work from many laboratories indicates that all three known selectins recognize sialylated fucosylated lactosaminyl oligosaccharides, and perhaps some sulfated glycoconjugates as well (2, 7-19). The structural motif Siaa2-3Galβ1-4(Fucα1-3)GlcNAc, called sialyl-Lewis\(^+\) (SLe\(^+\))\(^3\) and its isomer Siaa2-3Galβ1-3(Fucα1-4)GlcNAc, called sialyl-Lewis\(^*\), can be recognized by all three known selectins (12, 13, 17-19). Since SLe\(^+\) is found in abundant amounts on the surface of myeloid cells, it is natural to assume that it plays a major role in recognition of such cells by E- and P-selectin. However, even partial inhibition of selectin interactions by the SLe\(^+\) tetrasaccharide requires concentrations in the \(\mu\text{M}\) to \(\text{mM}\) range, and direct binding requires multivalent presentation in the form of immunobiotin complexes, neoglycoproteins, or glycolipids (12, 13, 17, 19, 21). Other evidence indicates that the detailed structure of the natural ligands for each selectin molecule may be distinct from one another. Thus, the binding specificities of E-selectin and P-selectin on neutrophils appears to be different by many criteria, and specific glycoprotein ligands have been identified for P- and L-selectin (16, 19, 22-26). Because of the role of selectins in the early phases of leukocyte emigration and thrombosis (27-30), and the hope of altering this process in pathological states (31-33), the detailed structural analysis of these natural ligands has become a major priority.

We have previously shown that neutrophils and HL-60 cells express a distinct sialoglycoprotein that binds with high affinity to P-selectin. This glycoprotein (apparent molecular mass...
using a P-selectin blotting assay (26). It was found only in myeloid cells, could bind selectively to columns of immobilized P-selectin in a calcium-dependent manner, and could be metabolically labeled with [3H]glucosamine in HL-60 cells, which are of myeloid origin (26). Evidence was also presented that this protein was distinct from several known neutrophil sialoglycoproteins, such as lamp-1, lamp-2, L-selectin, or leukosialin (26). The sialic acid(s) on this novel ligand were required for binding by P-selectin but appeared to be somewhat resistant to sialidase treatment. We have therefore explored whether this resistance is due to sialic acid modification (e.g. with an O-acetate group) or to a specific linkage of the sialic acid to the underlying oligosaccharide structure. We have also explored the role of the side chain of the sialic acid in recognition and found a simple way to introduce a tritium label into this minor component of neutrophil membranes. Finally, we demonstrate the sensitivity of the P-selectin ligand to a recently described enzyme of unusual specificity called O-sialoglycoprotease (34, 35), and we have used this enzyme to explore the role of O-linked oligosaccharides in mediating P-selectin binding to intact cells.

EXPERIMENTAL PROCEDURES

Enzymes—New Castle disease virus sialidase was prepared as previously described (36), Vibrio cholerae sialidase and Pronase were from Boehringer Mannheim, Antrobacter ureaeacieni sialidase from Calbiochem Behring Corp. (LaJolla, CA), and peptide N-glycosidase F from Genzyme (Cambridge, MA). The O-sialoglycoprotease from Pasteurella hemolytica was a generous gift from Dr. Alan Mellors, University of Guelph, Canada. Under the conditions used, it is known to be free of other proteases or sialidases. The specific activity of the enzyme preparation is defined by glycoinophorin A as a substrate (1 μl cleaves 5 μg/h).

Radioisotopes—[3H]Glucosamine and [3H]sodium borohydride were from Du Pont New England Nuclear, and carrier-free Na218O from Amersham Corp.

Antibodies—The monoclonal antibody (mAb) CSLEX1 (37) against SLE was kindly provided by Dr. Terasaki, UCLA; Vectastain ABC kits and biotinylated human anti-p chain were from Vector Laboratories (San Jose, CA). HLe-1 is an "isoform-unrestricted CD45 antibody" or de-0-acetylation (40, 41), and changes in peak area were monitored to confirm the identification of different sialic acids.

HPLC Analysis of Released Sialic Acids—Sialic acids were released from samples by incubation with 5-10 milliunits of A. ureafaciens sialidase in 100 μl of 100 mM sodium acetate, pH 5.5, containing 0.5% Triton X-100 for 14-16 h at 37 °C under a toluene atmosphere, and 2 mM periodate in phosphate-buffered saline, pH 7.0, overnight at 4 °C. The pH was adjusted to 8.0 by addition of 1 mNaOH and [3H]NaBH4 (6-10 μCi) was added. After 1 h at room temperature, excess NaBH4 was added to complete the reduction. In an additional 1 h, the sample was quenched with acetic acid and the sample was separated from the unreacted [3H]NaBH4 by-products by gel filtration on a Bio-Gel P-2 column run in Buffer A (see "P-selectin Affinity Chromatography").

P-selectin Ligand—Samples were subjected to electrophoresis on 7.5% SDS-polyacrylamide gels under reducing conditions and transferred to Immobilon-P membranes. The membranes were blocked and probed with 125I-labeled P-selectin (0.5-1 nM) as previously described (26). After extensive washing, they were exposed to X-Omat AR film at -70 °C for 1-10 h.

Release and Purification of Sialic Acids—Sialic acids were released from samples by incubation with 5-10 milliunits of A. ureafaciens sialidase in 100 μl of 100 mM sodium acetate, pH 5.5, containing 0.5% Triton X-100 for 14-16 h at 37 °C under a toluene atmosphere. Released sialic acids were purified by dialysis, followed by sequential ion exchange chromatography on Bio-Rad AG50 1×8 (hydrogen form) and Bio-Rad AG3×4A (formate form) as previously described (40, 41). These purification conditions were previously shown to avoid destruction of O-acetyl esters and to limit migration of 7-O-acetyl esters to less than 10%.

HPLC Analysis of Released Sialic Acids—Sialic acid-released sialic acids were purified, derivatized with DMB, and analyzed by reverse-phase HPLC as described (41, 42), using a TSK-gel ODS-120T column with an isotropic run of CH3CN:MeOH:H2O (9:7:84) at a 0.9 ml/min flow. Fluorescence was monitored with a SpectroVision Fluorometer using excitation and emission settings of 373 and 448 nm, respectively. Derivatization was done before and after induced migration or de-O-acetylation (40, 41), and changes in peak area were monitored to confirm the identification of different sialic acids.

HPLC Analysis of Underivatized Sialic Acids—The P-selectin ligand was metabolically labeled with [3H]glucosamine in HL-60 with as described above and affinity-purified. This purified ligand was fractionated by SDS-PAGE under reducing conditions and the 120-kDa ligand was excised and digested with Pronase. The resulting labeled glycopeptides were then digested with A. ureafaciens sialidase as described above and the released sialic acids separated from the unreacted material by gel filtration on a Bio-Rad P-2 column in 100 mM ammonium acetate. The released sialic acids were analyzed on a Varian AX-5 HPLC column under isocratic conditions of CH3CN:

2 A. Mellors, personal communication.
H$_2$O(0.25 M)NaHPO$_4$, pH 4.45 (72:18:10), at 1 ml/min (41). Fractions of 0.33 min were collected for scintillation counting.

**Jacalin Lectin Binding**—Samples were applied to a 2-ml Jacalin-Sepharose column (2 mg lectin/ml resin) equilibrated with 0.1 M NaCl, 10 mM MOPS, pH 7.5, 2 mM CaCl$_2$, 2 mM MgCl$_2$, 0.1% Triton X-100. The column was washed with equilibration buffer and eluted with 100 mM α-methylgalactoside in the same buffer. One-ml fractions were collected directly into scintillation vials and counted.

β-Elimination—The [3H]glucosamine-labeled or [3H]sialic acid-labeled P-selectin ligands were β-eliminated as described previously (43, 44). Following the alkaline-borohydride treatment, the samples were digested with Pronase (10 mg/ml) overnight at 60 °C. The samples were boiled and then passed over a 3-ml column of Dowex-50 (H$^+$ form) to remove Na$^+$ and amino acids. The unbound radioactivity was then analyzed by chromatography on Bio-Gel P-10 as described previously (45).

**O-Sialoglycoprotease Treatments**—Samples of the P-selectin ligand were incubated with 1 μg of enzyme in 100 mM HEPES, pH 7.4, in the presence of bovine serum albumin (stabilizer) for 1 h at 37 °C. The enzyme was inactivated by heating at 100 °C for 5 min. Samples (50 μg) of WGA eluate were incubated with or without 1 μg O-sialoglycoprotease in 0.1 M NaCl, 20 mM MOPS, pH 7.5, 0.02% NaN$_3$, for 18 h at 37 °C. HL-60 cells (100 μl, 10$^5$ cells/ml) in HBSS, 1% FCS, 0.1% NaN$_3$, 10 mM Neutrogen (sialidase inhibitor) were incubated for 60 min at 37 °C in the presence or absence of 2 μl of O-sialoglycoprotease. The cells were then washed with HBSS, 1% FCS, 0.1% NaN$_3$ prior to immunostaining.

**SDS-PAGE/Fluorography**—Samples were pooled, lyophilized, and boiled in sample buffer with or without 2-mercaptoethanol. Gels were run at 100 V constant until dye fronts ran to the bottom of the gel, stained with Coomassie Blue, and then exposed to En3Hance as per the manufacturer's directions. The dried gels were then exposed to X-Omat AR film at −70 °C and developed.

**Flow Cytometry**—P-selectin binding to HL-60 cells was assessed as previously described (46). Briefly, 10$^6$ HL-60 cells were incubated with 50 μl of purified P-selectin (10 μg/ml) in the presence or absence of 20 μg/ml G1, a mAb which inhibits P-selectin binding to myeloid cells. Bound P-selectin was detected by sequential incubation of cells with biotin-conjugated S12 (10 μg/ml) followed by 20 μg of phycoerythrin-streptavidin (undiluted). For immunostaining with Leu-22 (anti-CD43), HLe-1 (anti-CD45), and CSLEX1 (anti-sialyl-Le$^a$), 10$^6$ HL-60 cells were incubated with 2 μg/ml mAb followed by 50 μl of FITC-conjugated goat anti-mouse IgG/IgM (25 μg/ml). Each incubation was for 30 min at 4 °C in HBSS, 1.0% FCS, 0.1% NaN$_3$, between which the cells were washed with HBSS, 1.0% FCS, 0.1% NaN$_3$. After the last wash the cells were fixed with 1% paraformaldehyde and analyzed using a Becton-Dickinson FACScan flow cytometer.

**HL-60 Cell Adhesion Assay**—P-selectin was coated onto microtiter plates (5 μg/ml in HBSS, 100 μl/well) overnight at 4 °C. The wells were washed twice with HBSS and blocked with 300 μl of 0.1% casein in HBSS for 2 h at 22 °C. After washing the wells three times with HBSS, HL-60 cells were added (2 x 10$^4$/ml in HBSS, 1% FCS, 100 μl/well) and incubated for 20 min at 22 °C. The cells were filled with HBSS, sealed with acetate tape, and inverted for 10 min. Non-adherent cells were removed and the number of adherent cells was quantified by myeloperoxidase activity as previously described (38). All assays were performed in triplicate.

**RESULTS**

A 120-kDa P-selectin Ligand on Neutrophils Carries α2-3-Linked Sialic Acids That Are Relatively Resistant to Sialidases—In a previous study, we noted that binding of 125$I$-labeled P-selectin to a 120-kDa ligand from neutrophils and HL-60 cells was abolished by prolonged digestion of WGA-enriched sialoglycoproteins with sialidase from A. ureafaciens, which can release both α2-6- and α2-3-linked sialic acids (26). However, short digestions resulted only in an increase in apparent $M_r$ of the P-selectin ligand, indicating that substantial removal of sialic acids had occurred without much loss of binding. These observations suggested that there might be two or more populations of sialic acids that differ with respect to their linkage to the underlying oligosaccharide and/or the presence of modifications such as O-acetyl esters. To study the linkage of the sialic acids, we treated the WGA-enriched glycoproteins from neutrophil membranes with the sialidase from New Castle disease virus (NDV), which specifically cleaves α2-3-linked sialic acids (whether or not they are O-acetylated), or sialidase from V. cholerae, which cleaves α2-6- and α2-3-linked sialic acids but is less tolerant of sialic acids with modifications (47, 48). As shown in Fig. 1, a short treatment with these enzymes caused an upward shift in the apparent $M_r$ of the P-selectin ligand without affecting P-selectin binding. The effect of V. cholerae sialidase treatment was very similar to that obtained with the enzyme from Arthrobacter (data not shown). NDV treatment caused some upward shift in apparent $M_r$, indicating that α2-3-linked sialic acids are present on the P-selectin ligand. Although the shift with NDV was not as great as that seen with either A. ureafaciens or V. cholerae sialidase, prolonged treatment with the NDV sialidase was not attempted since the enzyme is not very stable to longer incubations and its specificity is not as stringent under such conditions. Thus, while α2-3-linked sialic acids are present on the P-selectin ligand, we cannot rule out the possibility that there are also sialic acids in α2-6 linkage. Regardless, the fact that some sialic acids on the ligand are relatively resistant to several sialidases raises the possibility that they might be substituted (49).

**Mild Periodate Oxidation of the Sialic Acid Side Chain Does Not Abolish Recognition of the Ligand by P-selectin**—Mild periodate oxidation (2 mM at 4 °C) can be used to selectively cleave the side chains of sialic acids while maintaining the integrity of the underlying oligosaccharide and protein struc-

---

3. B. K. Hayes and A. Varki, unpublished observations.
N-Acetylenuraminic Acid is the Single Major Sialic Acid on the P-selectin Ligand from Neutrophils and from HL-60 Cells—Since two lines of evidence suggested the possibility of sialic acid substitutions, it was necessary to examine this question directly. The sialidase from A. ureafaciens has a broad spectrum of action against most, but not all, sialic acids in a variety of types and linkages. However, since recognition of the 120-kDa ligand by P-selectin was completely abolished by the Arthrobacter enzyme, it is reasonable to assume that the sialic acids relevant for binding are released by this enzyme. Sialic acids from total neutrophil membranes were released with A. ureafaciens sialidase and compared with those from a WGA-enriched pool and from the P-selectin ligand purified by affinity chromatography on a P-selectin column. The released sialic acids were purified, derivatized, and analyzed by a TSK-ODS HPLC system that has previously been shown to detect most major forms of substituted sialic acids (41, 42). As shown in Fig. 2, the major sialic acid in all fractions studied was N-acetylenuraminic acid (Neu5Ac), the unmodified parent sialic acid. Although a very small amount of 9-O-acetyl-Neu5Ac was seen in the starting material, no enrichment of this species was seen in the purified ligand.

The P-selectin ligand isolated from neutrophils shares many properties with a similar protein found on HL-60 cells (26). To study the sialic acids on the HL-60 cell ligand, these cells were labeled with [6-3H]glucosamine. Because of metabolic interconversions, the tritium from [6-3H]glucosamine will be found in GlcNAc, GalNAc, and sialic acid residues (56). This makes it possible to enzymatically release, purify, and analyze radiolabeled sialic acids from a glycoprotein. The P-selectin ligand isolated from HL-60 cells labeled with [6-3H]glucosamine was purified by affinity chromatography on a P-selectin column. After separation by SDS-PAGE under reducing conditions, the radioactivity in the P-selectin ligand band was released from the gel by Pronase digestion as described under "Experimental Procedures." Released glycopeptides were large enough to void the Bio-Gel P-2 column. Treatment with A. ureafaciens sialidase released ~30% of the label as free sialic acids, which were separated from the remaining glycopeptides by Bio-Gel P-2 gel filtration. The released sialic acids were purified and analyzed using an AX-5 HPLC system, which can separate most major forms of substituted sialic acids but requires no derivatization prior to analysis. As shown in Fig. 3, we only found evidence for the unmodified sialic acid, [3H]Neu5Ac. These data indicate that the sialic acids responsible for P-selectin binding to the 120-kDa ligand are not substituted in neutrophils or in HL-60 cells. Rather, the partial resistance to sialidase treatment might be determined by features such as steric hindrance by the polypeptide and/or close spacing of the oligosaccharides.

The P-selectin Ligand Carries a Very Small Fraction of the Total Glycoprotein-bound Sialic Acids of the Neutrophil—Using the sensitive fluorescent HPLC assay, the yield of sialic acids was monitored in a purification in which quantitative recovery of the P-selectin ligand was obtained at every step as determined by P-selectin blotting (data not shown). Starting from a membrane extract containing 852 nmol of sialic acids, the final preparation of ligand contained only 0.74 nmol (0.09%) of the starting sialic acids. Thus, the amount of material available for direct analysis was very small. We took advantage of the fact that the ligand carries unmodified Neu5Ac whose side chains are not required for P-selectin binding, and specifically introduced tritium label into the sialic acid by truncation of the side chain with mild periodate oxidation, followed by reduction with NaBH₄ (50). A WGA-enriched glycoprotein fraction from neutrophils was first labeled in this manner, resulting in the introduction of tritium into many sialoglycoproteins. When applied to a P-selectin affinity column, almost all of the radioactivity ran through. However, a minor fraction of the tritium was specifically eluted with 5 mM EDTA (see Fig. 4, upper panel). When this small peak (representing 0.29% of the total radioactivity loaded on the column) was pooled, concentrated by acetone precipitation, and studied by SDS-PAGE and fluorography, a single glycoprotein that coincides in apparent molecular...
weight with the P-selectin ligand was seen after prolonged exposure (Fig. 4, lower panel). Note that this band migrated slightly faster than the prominent high molecular weight band found in the unbound material, which probably represents leukosialin (CD43), the major sialoglycoprotein of neutrophils (20).

The P-selectin Ligand Carries a Small Fraction of the Sialyl-Lewis\(^x\) Sequences on Neutrophil Membranes—\(\alpha\)2-3-Linked Neu5Ac is very commonly found at the non-reducing termini of oligosaccharide structures. It is also the terminal component of the less common tetrasaccharide motif Siao2-3Gal\(\beta\)1-4(Fuc\(\alpha\)1-3)GlcNAc (SLe\(^x\)), which has been demonstrated to be a basic recognition sequence for P-selectin (12, 13, 16, 17, 25). We therefore probed the myeloid P-selectin ligand for the presence of SLe\(^x\) using CSLEX1, an antibody that specifically recognizes this tetrasaccharide at the end of an oligosaccharide chain (37, 57). As shown in Fig. 5, all of the CSLEX1-positive material from a neutrophil membrane lysate bound to a WGA column, and the 120-kDa ligand purified from the P-selectin column eluate was also recognized by CSLEX1. However, the great majority of the CSLEX1-positive glycoproteins did not bind to the P-selectin column, indicating that the P-selectin ligand carries only a small fraction of the total SLe\(^x\) sequences in the neutrophil membrane. Thus, the SLe\(^x\) motif is not by itself sufficient to explain the high affinity binding of P-selectin to the 120-kDa ligand. Taken together with the data in the preceding section, these results indicate that the P-selectin ligand is a very minor sialoglycoprotein on the neutrophil. This observation corroborates well with previous data indicating a small number of high affinity binding sites for P-selectin on neutrophils and HL-60 cells (16, 22).

Jactalin Lectin Binding and \(\beta\)-Elimination Indicate the Presence of O-Linked Oligosaccharides on the P-selectin Ligand—Our previous studies indicated that, whereas the ligand appeared to be extensively sialylated (as evidenced by the upward shift in apparent molecular weight upon sialidase treatment), N-linked oligosaccharides constituted only a minor portion of its molecular weight, suggesting that most of the sialic acid might be on O-linked structures (26). The very small amount of material available made it difficult to carry out direct structural analysis of the oligosaccharides on this protein. We therefore sought alternate evidence concerning the nature of these sugar chains and tested for the ability of these oligosaccharides to interact with the Jactalin lectin, which recognizes the core of O-linked oligosaccharides (58, 59). We found that the majority of the radioactivity in the \([\text{H}]\)glucosamine-labeled ligand purified from HL-60 cells bound to a column of immobilized Jactalin lectin, suggesting that it does carry O-linked oligosaccharides (data not shown).

To obtain additional evidence for the presence of O-linked oligosaccharides on the P-selectin ligand, we subjected the \([\text{H}]\)glucosamine-labeled protein to \(\beta\)-elimination, a procedure that releases O-linked oligosaccharides from the peptide backbone (see “Experimental Procedures”). As shown in Fig. 6, most of the label from both the \([\text{H}]\)glucosamine-labeled and the periodate-[\(\text{H}\)]borohydride-labeled P-selectin ligand was...
The amounts of protein loaded onto the lanes were as follows: 200 pg of membrane extract and WGA flow-through electrophoresis on eluate and P-selectin flow-through, and 1 pg of P-selectin eluate. The carried on O-linked sugar chains. Since our prior work suggested the presence of only one or two N-linked chains that CSLEX1. Bound antibody was detected using biotin anti-p chain-same membrane was then re-probed with the anti-SW mAb from the indicated steps of the isolation procedure were subjected to Ion membranes, and probed with '251-labeled P-selectin specific antibody followed by streptavidin horseradish peroxidase released as oligosaccharides that eluted in the included volume on a Bio-Gel P-10 column. This indicates that most of the sialic acids and hexosamines on the P-selectin ligand are therefore re-examined the question of substrate specificity of the enzyme on the P-selectin ligand affinity-purified from [3H] glucosamine-labeled HL-60 cells, and on the detection of the ligand by P-selectin blotting of neutrophil membrane proteins. In both cases, the P-selectin ligand was also destroyed by O-sialoglycoprotease (Fig. 7, B and C). These results indicate that the P-selectin ligand has features that are specifically recognized by the O-sialoglycoprotease.

The O-Sialoglycoprotease from P. hemolytica Selectively Cleaves Glycoproteins with Many Closely Spaced Sialylated O-Linked Oligosaccharides—Previous reports concerning the substrate specificity of this enzyme indicate that it recognizes and cleaves only glycoproteins that carry sialylated O-linked oligosaccharides. However, as seen in Fig. 7A, the O-linked sialoglycoprotein fetuin was not affected by this enzyme. We therefore re-examined the question of substrate specificity of O-sialoglycoprotease, by reviewing the literature and by direct analysis of cleavage of selected proteins by the enzyme. We found that O-sialoglycoprotein cleaved bovine submaxillary mucin, κ-casein, and the recently described L-selectin ligand found in supernatants of lymph nodes (24), but that it did not cleave plasminogen (data not shown). From the information compiled in Table I and from other data in the literature, the following observations can be made. First, proteins without sugar chains and glycoproteins with only N-linked oligosaccharides are not cleaved. There-
was directly detected by transferring to Immobilon membranes and probing with $^3$H-labeled P-selectin.

Radiolabeled by periodate oxidation/$^3$HNaBH$_4$ reduction and detected by fluorography; C, P-selectin ligand in a WGA eluate from a neutrophil membrane lysate and analyzed by SDS-PAGE under reducing conditions.

<table>
<thead>
<tr>
<th>Glycoprotein</th>
<th>Total O-linked sites</th>
<th>O-Sialoglycoprotease sensitivity</th>
<th>Occurrence of linear stretches of amino acids with different numbers of adjacent O-linked sites$^a$</th>
</tr>
</thead>
<tbody>
<tr>
<td>Immunoglobulin A1 (44, 64)</td>
<td>1</td>
<td>Resistant (35)</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>Transferrin receptor (62, 63)</td>
<td>1</td>
<td>Resistant (34)</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>Fetuin (61)</td>
<td>3</td>
<td>Resistant$^b$</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>Plasminogen (60)</td>
<td>1</td>
<td>Resistant$^b$</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>Glycophorin A (108)</td>
<td>15</td>
<td>Sensitive (35)</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>Sgp50 L-selectin ligand from lymph nodes (24)</td>
<td>42$^c$</td>
<td>Sensitive$^{a,b}$</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>Bovine $\alpha$-casein (109)$^d$</td>
<td>31$^e$</td>
<td>Sensitive$^a$</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>Bovine submaxillary mucin (110)</td>
<td>163$^f$</td>
<td>Sensitive$^a$</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>CD43 (leukosialin) (111)</td>
<td>112$^g$</td>
<td>Sensitive (34)</td>
<td>1 2 3 4 &gt;4</td>
</tr>
<tr>
<td>CD34 (112)</td>
<td>95$^h$</td>
<td>Sensitive (34)</td>
<td>1 2 3 4 &gt;4</td>
</tr>
</tbody>
</table>

$^a$This refers to the spacing of O-linked glycosylation sites along the linear protein backbone, e.g. for glycophorin A, there are six individual attachment sites, one group of three adjacent sites (Ser$_2$, Thr$_2$, Thr$_3$), and one group of six adjacent sites (Thr$_{10}$, Thr$_{11}$, Thr$_{12}$, Ser$_{13}$, Ser$_{14}$, Ser$_{15}$).

$^b$Proteins examined in this study. The individual proteins were incubated with 1 ml of the enzyme at 37°C for 1 h and re-examined by SDS-PAGE and Coomassie Blue staining, PAS staining, or autoradiography as appropriate.

$^c$In these instances, all of the glycosylation sites have not been precisely identified. These numbers represent the total number of serine and threonine residues in the protein, and therefore the maximum number of O-linked sites possible. In these “mucin-type” proteins, most but not all of these sites are O-glycosylated.

$^d$Independently noted to be sensitive by Dr. A. Mellors (personal communication). Note: thus far, serum glycoproteins with N-linked oligosaccharides (34, 35), and many other proteins and glycoproteins that contain no O-linked oligosaccharides (34, 35) have proven to be resistant to O-sialoglycoprotease. Additionally, CD45 isoforms which show extensive but variable O-glycosylation in the amino-terminal domains show variable sensitivity to O-sialoglycoprotease.

Oligosaccharides. Since the P-selectin ligand from myeloid cells is cleaved by this enzyme, it is predicted to be a mucin-type glycoprotein.

**O-Sialoglycoprotease Treatment of HL-60 Cells Eliminates Binding to P-selectin.**—The data presented in this and a previous paper (26) indicate that the 120-kDa ligand binds P-selectin with high affinity. However, they do not directly demonstrate that this ligand accounts for the small number of high affinity binding sites observed on intact neutrophils or HL-60 cells. The O-sialoglycoprotease will not cleave cell surface glycoproteins with N-linked oligosaccharides or cell surface glycolipids (34, 35). On the other hand, it selectively cleaves cell-surface mucin-like molecules such as CD34 from intact cells (34, 65) and allows subsequent recovery of viable stem cells (66). We therefore examined the effects of this enzyme upon the direct binding of P-selectin to HL-60 cells. As seen in Fig. 8 (panels A and B), HL-60 cells were tested for their ability to bind soluble P-selectin after treatment with O-sialoglycoprotease. Sham-treated cells maintained their ability to bind to P-selectin, and this binding was blocked by co-incubation with the mAb G1, which has been previously shown to block P-selectin binding to myeloid cells (22, 38, 39). Treatment with O-sialoglycoprotease abolished binding by P-selectin. As controls, we also examined the effects of O-sialoglycoprotease on other sialylated surface molecules on HL-60 cells. As seen in Fig. 8 (panels C and D), binding of a mAb to CD43 (leukosialin) was abolished by O-sialoglycoprotease treatment, whereas binding of HL-E-1, an isoform-unrestricted CD45 mAb was unaffected. These results are consistent with previous work, which indicates that the...
either sham-treated or treated with lectin of detected with biotin-S12 and phycoerythrin-streptavidin as described under "Experimental Procedures." The washed cells (10⁶) were assayed for direct binding of P-selectin in the presence or absence of 20 μg/ml of the blocking mAb G1 (panels A and B). Bound P-selectin was detected with biotin-S12 and phycoerythrin-streptavidin as described under "Experimental Procedures." Cells were also analyzed for surface expression of CD43 (Leu-22), CD45 (HLe-1), and sialyl-Le¹ (CSLEX1) by incubation with 2 μg/ml indicated mAb followed by FITC-conjugated goat anti-mouse IgG/IgM F(ab')₂ (panels C and D).

Our previous work has suggested a role for a specific cell-surface glycoprotein ligand in the high affinity recognition of neutrophils and HL-60 cells by P-selectin (26). In this study, we have looked more closely at the nature of the sialic acids and the oligosaccharides involved in this binding. Using sialidases with linkage specificity, we have shown that at least some of the sialic acids on the P-selectin ligand are in α2-3 linkage to the underlying oligosaccharides. While it is possible that all of the sialic acids are in this linkage, we could not address this issue, because the NDV sialidase used to demonstrate α2-3 linkages can also cleave α2-6 linkages with prolonged exposure. Thus, we cannot rule out the presence of α2-6-linked sialic acids on the ligand (25). An even less likely possibility is the presence of α2-8-linked sialic acids that can be cleaved by both A. ureafaciens sialidase and NDV; however, this type of linkage has been reported on only a few specific glycoproteins in nature (67). The relative resistance of the sialic acid residues on the P-selectin ligand to several sialidases and the lack of effect of mild periodate oxidation on binding raised the suspicion that the relevant sialic acid residues were naturally modified (49). However, direct analysis indicates that the common unmodified sialic acid, Neu5Ac, is the predominant sialic acid form carried by the ligand from both neutrophils and HL-60 cells. Thus, the relative resistance to sialidase is probably due to steric factors such as close-spacing of oligosaccharides and/or protein conformation.

**FIG. 8. Effect of O-sialoglycoprotease on the direct binding of P-selectin to HL-60 cells.** HL-60 cells were either sham-treated or treated with O-sialoglycoprotease as described under "Experimental Procedures." The washed cells (10⁶) were assayed for direct binding of P-selectin in the presence or absence of 20 μg/ml of the blocking mAb G1 (panels A and B). Bound P-selectin was detected with biotin-S12 and phycoerythrin-streptavidin as described under "Experimental Procedures." Cells were also analyzed for surface expression of CD43 (Leu-22), CD45 (HLe-1), and sialyl-Le¹ (CSLEX1) by incubation with 2 μg/ml indicated mAb followed by FITC-conjugated goat anti-mouse IgG/IgM F(ab')₂ (panels C and D).

**FIG. 9. Effect of O-sialoglycoprotease on adhesion of HL-60 cells to immobilized P-selectin.** HL-60 cells were sham-treated or treated with active or heat-inactivated O-sialoglycoprotease (O-SGP). Adhesion of the cells was then quantified on microtiter wells coated with or without P-selectin and then blocked with casein. Adhesion is presented as the mean ± S.D. of triplicate determinations. Similar results were obtained in a second independent experiment. Adhesion of untreated cells (not shown) and sham-treated cells was similar. Flow cytometric analysis of the sham- and enzyme-treated cells used in this experiment demonstrated results identical to those shown in Fig. 8.
In this study, we have also shown that the side chain of sialic acids is not important for the binding of P-selectin to the ligand, and we have taken advantage of this fact to introduce high specific activity tritium label into this minor sialoglycoprotein. Other studies have indicated that the side chain of sialic acids is not required for recognition by E-selectin (14) or L-selectin (68). This observation contrasts with the absolute requirement for the side chain in recognition of ligands by sialoadhesin (69), influenza virus hemagglutinins (70), CD22 (71, 72), and certain monoclonal antibodies (53, 73). It remains to be seen what, if any effects, the natural modifications of side chains (e.g., O-acetylation) will have on recognition of sialylated ligands by the selectins.

A large body of prior evidence indicates that binding by all three selectins can involve either $\text{SLe}^\alpha$ (sialyl-Lewis$^\alpha$) or $\text{SLe}^\beta$ (sialyl-Lewis$^\beta$) as a minimal recognition structure (2, 7–19). We have shown that the P-selectin ligand contains the $\text{SLe}^\alpha$ sequence that is recognized by the CSLEX1 antibody. However, the great majority of the glycoprotein-bound $\text{SLe}^\alpha$ from neutrophils failed to bind to P-selectin with similar affinity. This correlates well with the finding that the P-selectin ligand carries a very small fraction of the total sialic acids from neutrophil or HL-60 cells. Taken together with other recent reports (7, 12, 16, 26), our present study indicates that if $\text{SLe}^\alpha$ is necessary for binding, it is not sufficient for high affinity binding of this myeloid ligand by P-selectin. In this regard, it is striking that treatment of HL-60 cells with O-sialoglycoprotease destroys P-selectin binding, but leaves total cell surface $\text{SLe}^\alpha$ largely intact (see Fig. 8). These data also indicate that other newly proposed ligands for P-selectin such as sulfatide (15, 74, 75) and sulfated glycosaminoglycans (76, 77) cannot be responsible for high affinity P-selectin binding, at least on myeloid cells. The paucity of the P-selectin ligand on myeloid cells also fits well with the small number of high affinity binding sites for P-selectin on these cells (22, 76). There is as yet no direct evidence that these sites can be directly attributed to, and completely accounted for, by this particular P-selectin ligand. However, our finding that O-sialoglycoprotease treatment of HL-60 cells destroys binding to P-selectin supports this notion.

The recognition of $\text{SLe}^\alpha$ by the selectins can be compared to some other situations involving specific biological recognition. For example, the discovery of the critical motifs involved in recognition of lysosomal enzymes by certain trafficking receptors (mannose 6-phosphate) or of matrix proteins by the integrins (RGD sequences) were landmark events (78, 79). However, basic recognition motifs have relatively low affinities for their receptors. Subsequent studies showed much greater complexity in the natural ligands for these receptors (80, 81), with biologically relevant recognition requiring other specific features of the ligands such as tertiary structure and/or multivalency. Thus, while sialylated fucosylated lactosamines such as $\text{SLe}^\alpha$ may form basic recognition motifs for the selectins, biologically relevant binding may require more complex structures such as those presented by the P-selectin ligand on myeloid cells described here. $\text{SLe}^\alpha$ found on other glycoconjugates in the neutrophil surface might also support cell adhesion if such low affinity interactions, due to their density, resulted in significant avidity on an intact cell surface. However, Chinese hamster ovary cells transfected with a fucosyltransferase, while expressing large amounts of $\text{SLe}^\alpha$ on their surface, lack high affinity binding sites for P-selectin and adhere only weakly to immobilized P-selectin (16). Here we demonstrate that HL-60 cells treated with O-sialoglycoprotease do not adhere to P-selectin, even though total surface expression of $\text{SLe}^\alpha$ is not detectably altered. The enzyme-treated HL-60 cells might bind to wells coated with higher densities of P-selectin like those used for the transfected Chinese hamster ovary cells. However, the weak adhesion of the latter cells may reflect the fact that they express much higher levels of $\text{SLe}^\alpha$ than do HL-60 cells (16). Therefore, low affinity ligands for P-selectin may not play a major role in adhesion unless perhaps they are positioned or clustered in favorable sites such as pseudopodia on the cell surface (82).

This study and our previous results (26) indicate that the P-selectin ligand has very few N-linked oligosaccharides and many O-linked sugar chains. O-Sialoglycoprotease is a recently described enzyme from $P.\ hemolytica$ that has been reported to cleave only sialylated O-linked glycoproteins (34, 35). Based on the glycoproteins that are reported substrates for this enzyme, we suspected that the specificity of this enzyme might be more complex than suggested. Taken together with the prior literature, the additional studies we have performed indicate that besides the presence of sialylated O-linked sugar chains, an additional requirement of the enzyme appears to be the presence of a large number oligosaccharides that are closely spaced and/or adjacent to one another. Of course, the exact recognition moiety may be only a subgroup of the total oligosaccharides present. Since the P-selectin ligand is susceptible to O-sialoglycoprotease, it is reasonable to suggest that it also carries closely spaced groupings of sialylated O-linked oligosaccharides. Such close-packing or "clustering" may also be the cause of the relative resistance to sialidase treatments.

The observation that the ligand can bind P-selectin even after denaturation with SDS and disulfide bond reduction suggests that P-selectin recognition is due solely to lectin-carbohydrate interaction. The specificity of P-selectin for the 120-kDa ligand might therefore be due to a unique structural feature of the oligosaccharide(s). For example, the protein backbone of the ligand might specify the addition of uncommon oligosaccharide sequences during biosynthesis, as is seen with mannose 6-phosphorylation of lysosomal enzymes (83, 84). However, it is also possible that the O-linked oligosaccharides on the 120-kDa ligand are not unusual or distinct from those on other neutrophil glycoproteins. In this case, the specificity of P-selectin for the ligand may be conferred by co-recognition of peptide sequences and/or by the clustering of common oligosaccharide structures. Detailed structural analysis of the oligosaccharide on the 120-kDa ligand will be required to distinguish these possibilities.

It is worth noting that a large number of monoclonal antibodies known to detect "tumor-specific" glycoproteins from normal and cancer cells have "sialidase-sensitive" or "sialic acid-dependent" epitopes (see Refs. 73 and 85–87 for examples). Many monoclonal antibodies apparently specific for the proteins CD24 (88, 89), CD45 (90–93), and CD43 (94, 95) also show sialidase sensitivity. In the case of CD43 (leukosialin), a heavily glycosylated O-linked glycoprotein from leukocytes (96), it is rare to find a specific mAb that does not require sialylation for recognition (97). In fact, expression cloning of a glycosyltransferase required the co-expression of a specific polypeptide, leukosialin, in the same cell type to permit recognition by a specific mAb (98). In many such cases, when the sialyl-oligosaccharides on such proteins are directly examined (20), they are not unusual in structure compared with those from other proteins of the same cells. Thus, the oligosaccharides cannot by themselves have generated the specificity seen by the antibodies. Many of the proteins with such sialidase-sensitive epitopes are either mucins with clustered O-linked oligosaccharides or have large
numbers of closely spaced N-linked oligosaccharides. In similar fashion, the reactivity of a mucin-specific lectin from Sambucus sieboldiana (99) appears to require clustering of O-linked oligosaccharides and cannot recognize fetuin. Likewise, high affinity recognition of O-linked oligosaccharides on glycoprotein A by a 175-kDa invasion receptor of the F. palaeum merozoite appears to require the clustering of the sugar chains (100).

Oligosaccharides in free solution or at single attachment sites have a great deal of freedom of mobility and show limited interactions with associated proteins (101). However, oligosaccharides closely packed together (as in mucins) can form a relatively rigid rodlike structure (95, 102) and presumably have less freedom of mobility. We hypothesize that such clustering of common oligosaccharides might serve to present uncommon epitopes by forming a unique “clustered saccharide patch,” in a manner somewhat analogous to the recognition of DNA by specific transcription factors (103). This type of epitope or recognition site would be generated by a combination of side chains and groups from multiple oligosaccharides that are packed close enough to prevent excessive mobility. Although the polypeptide would not itself be part of the recognition patch, it would be crucial for presenting the sugar chains in the correct arrangement. If the arrangement was disturbed, recognition would be lost, and individual sugar chains would be too little (if any) for the affinity for the receptor in question. This type of recognition patch would thus be distinct from that described for certain lectins in which “correct spacing” of terminal or internal sugar residues provides improved binding of certain multiantennary oligosaccharides or glycopeptides with widely spaced non-adjacent O-linked oligosaccharides (104–107). If antibodies, plant lectins, and parasite receptors can specifically recognize closely spaced clustered O-linked oligosaccharides, it seems possible that mammalian lectins such as the selectins could do the same. Thus, while further structural analyses of the O-linked oligosaccharides of the P-selectin ligand might well uncover unique structures required for recognition, it is possible that they may not. In the latter case, it will be even more important to elucidate the polypeptide chain sequence and the structure and sites of oligosaccharide attachment on the protein.


REFERENCES

P-selectin Ligand