

Commentary & View

Potential of N-glycan in cell adhesion and migration as either a positive or negative regulator

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Glycosylation is one of the most abundant posttranslational modification reactions, and nearly half of all known proteins in eukaryotes are glycosylated. In fact, changes in oligosaccharide structure (glycan) are associated with many physiological and pathological events, including cell adhesion, migration, cell growth, cell differentiation and tumor invasion. Glycosylation reactions are catalyzed by the action of glycosyltransferases, which add sugar chains to various complex carbohydrates such as glycoproteins, glycolipids and proteoglycans. Functional glycomics, which uses sugar remodeling by glycosyltransferases, is a promising tool for the characterization of glycan functions. Here, we will focus on the positive and negative regulation of biological functions of integrins by the remodeling of *N*-glycans with *N*-acetylglucosaminyltransferase III (GnT-III) and *N*-acetylglucosaminyltransferase V (GnT-V), which catalyze branched *N*-glycan formations, bisecting GlcNAc and β 1,6 GlcNAc, respectively. Typically, integrins are modified by GnT-III, which inhibits cell migration and cancer metastasis. In contrast, integrins modified by GnT-V promote cell migration and cancer invasion.

Protein glycosylation encompasses *N*-glycans, *O*-glycans and Glycosaminoglycans. *N*-glycans are linked to asparagine residues of proteins, which is a specific subset residing in the Asn-X-Ser/Thr motif, whereas *O*-glycans are attached to a subset of serines and threonines (Fig. 1).¹ An increasing body of evidence indicates that glycans in glycoproteins are involved in the regulation of cellular functions including cell-cell communication and signal transduction.^{2,3} In fact, most receptors on the cell surface are *N*-glycosylated—integrins and epithelial growth factor receptors; and transforming growth factor β receptors. Here, we focus mainly on the modification of *N*-glycans of integrin α 3 β 1 and α 5 β 1 to address the important roles of *N*-glycans in cell adhesion and migration.

Previous studies indicate that the presence of the appropriate oligosaccharide can modulate integrin activation. When human fibroblasts were cultured in the presence of 1-deoxymannojirimycin, an inhibitor

of α -mannosidase II, which prevents *N*-linked oligosaccharide processing, immature α 5 β 1 integrin appeared at the cell surface, and fibronectin (FN)-dependent adhesion was greatly reduced.⁴ In addition, the treatment of purified integrin α 5 β 1 with *N*-glycosidase F, which cleaves between the innermost GlcNAc and asparagine residues of *N*-glycans from *N*-linked glycoproteins, resulted in the blockage of α 5 β 1 binding to FN and the inherent association of both subunits,⁵ suggesting that *N*-glycosylation is essential for functional integrin α 5 β 1. The production of glycoprotein glycans is catalyzed by various glycosyltransferases. *N*-Acetylglucosaminyltransferase III (GnT-III) transfers *N*-acetylglucosamine (GlcNAc) from UDP-GlcNAc to a β 1, 4 mannose in *N*-glycans to form a “bisecting” GlcNAc linkage, as shown in Figure 2. Bisecting GlcNAc linkage is found in various hybrid and complex *N*-glycans. GnT-III is generally regarded as a key glycosyltransferase in *N*-glycan biosynthetic pathways. Introduction of a bisecting GlcNAc suppresses further processing and elongation of *N*-glycans catalyzed by *N*-acetylglucosaminyltransferase V (GnT-V), which is strongly associated with cancer metastasis, since GnT-V cannot utilize the bisected oligosaccharide as a substrate.⁶⁻⁸ It has also been reported that GnT-V activity and β 1, 6 branched *N*-glycan levels are increased in highly metastatic tumor cell lines.^{9,10} When NIH3T3 cells were transformed with the oncogenic Ras gene, cell spreading on FN was greatly enhanced due to an increase in β 1, 6 GlcNAc branched tri- and tetra-antennary oligosaccharides in α 5 β 1 integrins.⁹ Similarly, the characterization of *N*-glycans of integrin α 3 β 1 from non-metastatic and metastatic human melanoma cell lines showed that β 1, 6 GlcNAc branched structures were expressed at high levels in metastatic cells compared with non-metastatic cells.¹⁰ Cancer metastasis was consistently, and significantly, suppressed in GnT-V knockout mice.¹¹

To explore the possible mechanisms involved in increased β 1, six branched *N*-glycans on cancer cells, Guo et al. found that cell migration toward FN and invasion through the matrigel were both substantially stimulated in cells in which the expression of GnT-V was induced.¹² Increased branched sugar chains inhibited the clustering of integrin α 5 β 1 and the organization of F-actin into extended microfilaments in cells plated on FN-coated plates, which supports the hypothesis that the degree of adhesion of cells to their extracellular matrix (ECM) substrate is a critical factor in regulating the rate of cell migration, i.e., migration is maximal under conditions of intermediate levels of cell adhesion.¹³ Conversely, GnT-V null mouse

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embryonic fibroblasts (MEF) displayed enhanced cell adhesion to, and spreading on, FN-coated plates with the concomitant inhibition of cell migration. The restoration of GnT-V cDNA in the null MEF reversed these abnormal characteristics, indicating the direct involvement of *N*-glycosylation events in these phenotypic changes.

In contrast to GnT-V, the overexpression of GnT-III resulted in an inhibition of $\alpha 5 \beta 1$ integrin-mediated cell spreading and migration, and the phosphorylation of the focal adhesion kinase.¹⁴ The affinity of the binding of integrin $\alpha 5 \beta 1$ to FN was significantly reduced as a result of the introduction of a bisecting GlcNAc to the $\alpha 5$ subunit. In addition, overexpression of GnT-III in highly metastatic melanoma cells reduced $\beta 1$, six branching in cell-surface *N*-glycans and increased bisected *N*-glycans.¹⁵ Therefore, GnT-III has been proposed as an antagonistic of GnT-V, thereby contributing to the suppression of cancer metastasis. In fact, the opposing effects of GnT-III and GnT-V have been observed for the same target protein, integrin $\alpha 3 \beta 1$.¹⁶ GnT-V stimulates $\alpha 3 \beta 1$ integrin-mediated cell migration, while overexpression of GnT-III inhibits GnT-V-induced cell migration. The modification of the $\alpha 3$ subunit by GnT-III supersedes modification by GnT-V. As a result, GnT-III inhibits GnT-V-induced cell migration. These results strongly suggest that remodeling of glycosyltransferase-modified *N*-glycan structures either positively or negatively modulates cell adhesion and migration.

In addition, sialylation on the non-reducing terminus of *N*-glycans of $\alpha 5 \beta 1$ integrin plays an important role in cell adhesion. The increased sialylation of the $\beta 1$ integrin subunit was correlated with a decreased adhesiveness and metastatic potential.¹⁷⁻¹⁹ On the other hand, the enzymatic removal of $\alpha 2$, eight-linked oligosialic acids from the $\alpha 5$ integrin subunit inhibited cell adhesion to FN,²⁰ supporting the observation that the *N*-glycans of α and β integrin subunits play distinct roles in cell-ECM interactions.²¹ Collectively, these findings suggest that the interaction of integrin $\alpha 5 \beta 1$ with FN is dependent on *N*-glycosylation and the processing status of *N*-glycans.

Although alteration of the oligosaccharide portion on integrin $\alpha 5 \beta 1$ could affect cis- and trans-interactions caused by GnT-III, ST6GalI and GnT-V, as described above, the molecular mechanism remains unclear. Considering integrin $\alpha 5 \beta 1$ contains 26 potential *N*-linked glycosylation sites (14 in the α subunit and 12 in the β subunit), the determination of those crucial *N*-glycosylation sites for its biological function is, therefore, quite important for an understanding of the underlying mechanism. We sequentially mutated either one or a combination of asparagine residues in the putative *N*-glycosylation sites of glutamine residues, and found that *N*-glycosylation on the β -propeller domain of the $\alpha 5$ subunit (in particular sites number 3–5) is essential for its hetero-dimer formation and its biological functions such as cell spreading and cell migration, as well as for the proper folding of the $\alpha 5$ subunit.²² On the other hand, *N*-glycans on $\beta 1$ integrin also play important roles in the regulation of its biological functions^{23,24} (and our unpublished data). Very recently, we also found that GnT-III specifically modifies one of the important glycosylation sites, which results in functional regulation (unpublished data). We postulate that these important sites may participate in supramolecular complex formation on the cell surface, which controls intracellular signal transduction.

It also is worth noting that *N*-glycans regulate cell-ECM association as well as cell-cell adhesion. Overexpression of GnT-III slowed E-cadherin turnover, resulting in increased E-cadherin expression on the surface of B16 melanoma cells.²⁵ E-cadherin engagement at

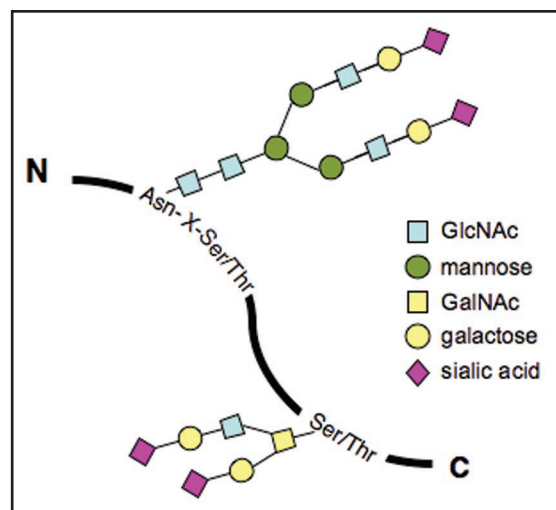


Figure 1. Two major types of protein glycosylation. *N*-glycans are covalently linked to asparagine (Asn) residue of proteins, specifically the Asn-X-Ser/Thr motif. In contrast, *O*-glycans are attached to a subset of glycosidically linked hydroxyl groups of the amino acids serine (Ser) and threonine (Thr).

cell-cell contacts is known to suppress cell migration, and that effect has been best described in the context of tumorigenesis.²⁶ Conversely, the disruption of E-cadherin-mediated cell adhesion appears to be a central event in the transition from non-invasive to invasive carcinomas. Interestingly, we recently found that E-cadherin-mediated cell-cell interaction upregulated GnT-III expression,^{27,28} suggesting that regulation of GnT-III and E-cadherin expression may exist as a positive feedback loop. Taken together, the overexpression of GnT-III inhibits cell migration by at least two mechanisms: an enhancement in cell-cell adhesion and a downregulation of cell-ECM adhesion (Fig. 2).

Indeed, glycosylation defects in humans and their links to disease have shown that the mammalian glycome contains a significant amount of biological information.²⁹ The mammalian glycome repertoire is estimated to be between hundreds and thousands of glycan structures and could be larger than its proteome counterpart. Nevertheless, characterization of the biological functions of each glycan could one day make a significant contribution to the diagnosis and treatment of disease.

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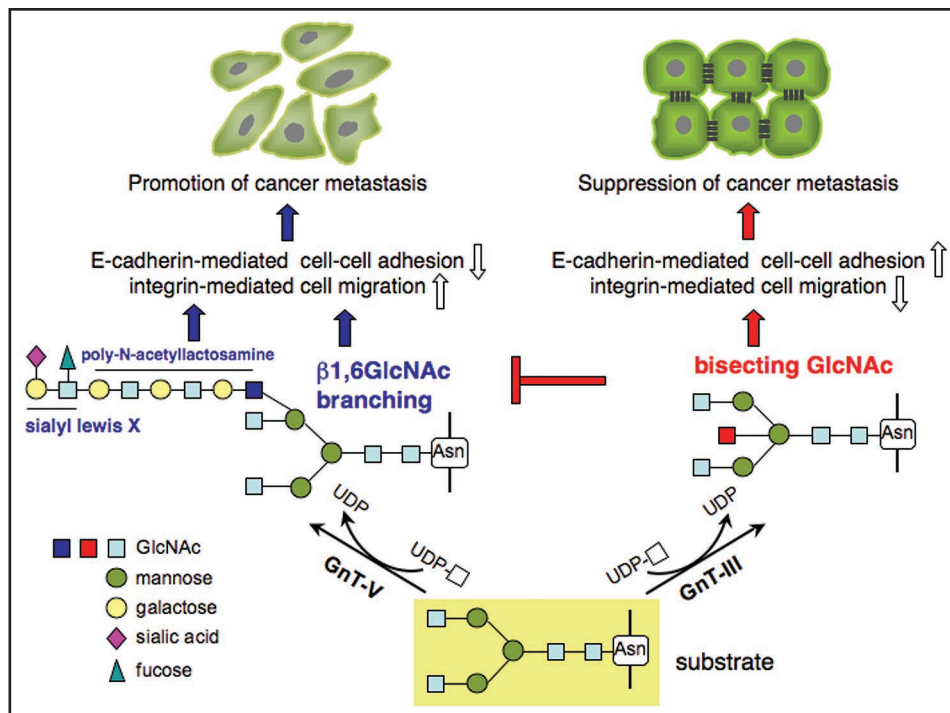


Figure 2. Glycosylation reactions catalyzed by the action of glycosyltransferase GnT-III and GnT-V. The remodeled N-glycans regulate cell adhesion and migration. Enhanced expression of GnT-V in epithelial cells results in a loss of cell-cell adhesion, increasing integrin-mediated cell migration. In contrast, overexpression of GnT-III strengthens cell-cell interaction and downregulates integrin-mediated cell migration, which may contribute to the suppression of cancer metastasis. The β 1,6GlcNAc branching is preferentially modified by polylactosamine and other sugar motifs such as sialyl Lewis X, which also contribute to promotion of cancer metastasis. It is worth mentioning that GnT-III could be proposed as an antagonistic of GnT-V, since GnT-V cannot utilize the bisected oligosaccharide as a substrate.

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